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Human neuropeptide
Human peptide YY (
Peptide YY SEQ ID
Human peptide tyro
Modified human pep
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Human peptide YY.
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Peptide Y. Syntho
Human peptide YY.
                                                                                                                        February 27, 2003, 14:15:13 ; Search time 35 Seconds
(without alignments)
137.058 Million cell updates/sec
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GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.
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Listing first 45 summaries
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AAR97741
AAW51801
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Gapop 10.0 , Gapext 0.5
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ALIGNMENTS

Peptide YY; PYY; gastro-enterological disorders; intestinal water; electrolyte secretion; cell proliferation; nutrient transport; lipolysis; blood flow regulation. New peptide derivs. - useful as therapeutic agents, for treating gastro-enterological disorders AAR62050 standard; peptide; 36 AA. 94WO-US03380 93US-0038534 93US-0109326 14-JUN-1995 (first entry) (UYCI-) UNIV CINCINNATI. Human peptide YY (PYY). WPI; 1994-332815/41. Balasubramaniam A; 29-MAR-1994; 29-MAR-1993; 19-AUG-1993; Homo sapiens WO9422467-A. 13-OCT-1994 AAR62050; RESULT 1 AAR62050 THE STATE OF THE S

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WPI; 1998-322327/28.
                                                                                         Sest Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36 AA;
                                                     36 AA;
                                                                                                                                                                                                                                                                                                  Human peptide YY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide YY; PYY; porcine; human; intestine; endocrine cell; gut motility; gastrointestinal tract; pancreas; inhibitor; intestinal secretion; pig; pancreatic tumour; blood flow; serous cyst adenoma; microcystic tumour; solid-cyst tumour; malignant tumour; therapy.
                                                                                                                                                                                                                                                 Gaps
                                 AAR62050 describes the amino acid sequence of human peptide YY (PYY), which was isolated from the endocrine cells of the human gastrointestinal tract and pencreas. Using the equivalent porcine PYY sequence (AAR62049) as a base the PYY analogues described in AAR62051-R62082 were produced. The new peptides were found to have a variety of properties, that made them useful as therapeutic agents in the treatment of gastro-enterological disorders. As part of a therapeutic composition they could be used for decreasing excess intestinal water and electrolyte secretion, for regulating regulating lipolysis and blood flow.
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                                                                                                                                                                                                                       Length 36;
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                                                                                                                                                                                                                    Score 194; DB 15;
Pred. No. 2.7e-20;
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100.0%; Pred. No. 2..
0; Mismatches
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             Disclosure; Page 3; 45pp; English
                                                                                                                                                                                                                                                                                                                                                              AAR97741 standard; peptide; 36 AA
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nes 36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide YY; cell proliferation; nutrient transport; lipolysis; electrolyte secretion; anti-secretory; intestinal water; antimotility.
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microcystic tumours, and solid-cyst tumours. The malignant tumours capable of being treated by the method of the invention include, carcinomas arising from the ducts, acini, or islets of the pancreas.
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Pred. No. 2.7e-20;
Mismatches 0;
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                                                                                                                                                                                                                         Pred. No. 2.7e-20;
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100.0%;
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us-09-634-363-2.rag

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NPY; NPY-Y4 receptor; precocious puberty; polycystic ovary syndrome; endometriosis; benign prostatic hyperplasia; delayed puberty; amenorrhea; breast cancer; prostate cancer;
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                                                                                                                                          Human; neuropeptide Y; luteinizing hormone; reproductive system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 2; 17pp; English.
                                                        20-JUN-2001 (first entry)
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                                                                                                    Human peptide YY
                                                                                                                                                                                                                           peptide YY; PYY
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                                                                                                                                                                                                                                                                                                                                  Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-NOV-1999;
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13-MAY-1999;
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                 AAB12178;
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This sequence represents peptide Y (PPY). The invention relates to neuropeptide (1a) of 3-10 amino acids (aa) in which at least two cyclic peptide (1a) of 3-10 amino acids (aa) in which at least two residues are joined by a naphthyl ring; and (ii) at least two related sequence PYY and PYY agonists, are used to reduce airway retaited sequence PYY and PYY agonists, are used to reduce airway restriction in patients with bronchists, are used to reduce airway restriction in patients may also be used: (i) to treat conditions bronchitis. The agonists may also be used: (i) to treat conditions responsive to NPY Or PYY, e.g. laryngitis, chronic rinnosinusitis, oedema, inflammation, anxiety, congestive heart failure, cardiomyopathy, coronary arterial disease, diminished cardiac vagal activity, hypertension, Alzheimer's disease, epilepsy, ischeemia, angina, myocardial infarction, acquired immune deficiency syndrome and diseases characterised by reduced immune responses; and (ii) to increase body weight or as an antihistamine. The template induces folding of (Ib) into a biologically active form. Since (1) contain only the C-terminal region of NPY, they are selective for the Y2 receptor.
                                                                                                                                                                                                                    Neuropeptide Y; NPY; agonist; peptide YY; PYY; airway restriction; bronchial disease; asthma: bronchiis; laryngitis; Alzheimer's disease; chronic rhinosinusitis; oedema: inflammation; congestive heart failure; cardiomyopathy; coronary arterial disease; myocardial infarction; AIDS; diminished cardiac vagal activity; hypertension; epilepsy; ischaemia; angina; immune response; antihistamine; therapy.
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to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New agonists of neuropeptide Y containing linear peptide linked cyclic template peptide, used e.g. to reduce airway restriction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 42; 45pp; English.
                                                      AAY43335 standard; peptide; 36 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Grouzmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-EP02076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0054393
                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lacroix J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BMRA-) BMRA CORP BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-620192/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                WO9951626-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-APR-1998;
                                                                                                                                       25-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-0CT-1999
                                                                                                                                                                                 Peptide Y.
                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mutter M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                AAY43335;
                            AAY43335
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Hayward A;

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Akinsanya

98GB-0025969. 99GB-0011178. 99WO-GB03963

/note= "C-terminal amide" Location/Qualifiers 36

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Neuropeptide Y (NPY) (AAB12177) has a number of effects on the reproductive system. NPY is one of a family of neuropeptides. Other members of the family include the present sequence, peptide YY (PYY), and pancreatic polypeptide (PP, See AAB1219) and AAB12180). Selective NPY-V4 receptor agonists have been found (see AAB12181 to AAB12183). The NPY-V4 receptor agonists cause an increase in the circulating levels of Luteinizing hormone (LH) and hence improve the fertility of animals with compromised reproductive disorders such as delayed puberty and amenorrhea. In addition, NPY-V4 antagonists may be used to treat human reproductive disorders such as precocious puberty, endometriosis, polycystic ovary syndrome, benign prostatic hyperplasia and hormonne-dependent neoplasias e. 9. breast cancer and prostate cancer. The present sequence was used in a sequence homology comparison.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 194; DB 21; 100.0%; Pred. No. 2.7e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                         36 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY87961;
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 6
AAY87961
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Gaps

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Indels

AAB12178 standard; peptide; 36

RESULT 5
AAB12178
ID AAB12

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The invention relates to novel peptide YY (PYY) analogues

(AAY87551-Y87564 and AAX87568) that can be used for the treatment of

CG (AAY87551-Y87564 and AAX87568) that can be used for the treatment of

CG gastrointestinal disorders. PYY (AAY87559, ARY87550) is a 36 residue

peptide amide that is released into the circulation after a meal and

CG which is thought to play a role in regulating intestinal secretion and

CC which is thought to play a role in regulating intestinal secretion and

CC inhibitor of diarrhoea. PYY has also been implicated in nutrient uptake,

CC call proliferation, I plotysis and vasoconstriction. The compounds of the

invention are useful for inhibiting fluid and electrolyte secretion in

CC call proliferation in the gastrointestinal tract; regulating lipolysis in,

CC example, adipose tissue; and regulating blood flow in mammals. The

canalogues are especially useful in the treatment of gastrointestinal

CC disorders associated with excess intestinal electrolytes and water

CC disorders associated with excess intestinal electrolytes and water

CC disorders associated with excess intestinal electrolytes and water

CC disorders associated with excess intestinal electrolytes and water

CC disorders associated with excess intestinal electrolytes and water

CC disorders associated with excess intestinal diarrhoea, diarrhoea due to

CC disorders alifertions, travellers' diarrhoea, inflammancory diarrhoea

CC or diarrhoea following lleostomy. The peptides can also be used to treat intestinal

CC diarrhoea following lleostomy. The peptides can also be used to treat intestinal

CC diarrhoea following lleostomy. The peptides can also de used to treat intestinal

CC diarrhoea following leostomy. The peptides can also be used to treat intestinal

CC diarrhoea following leostomy or useful the PAY analogues are shorter than naturally occurring PAY, suthers and purification of the compounds is easier and less costly. The

CC and purification of the compounds and plays of the PAY analogues interact spe
                                                                                                                                                                                                                                                                                                                                   New peptide YY analogs especially useful for treating gastrointestinal disorders associated with excess intestinal electrolytes and water secretion, and decreased absorption, e.g. infectious and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protection, endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     side reactions. The present sequence represents human PYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 194; DB 21;
ilarity 100.0%; Pred. No. 2.7e-20;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Column 2; 18pp; English.
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                                                                            980S-0047986
                                                                                                                            98US-0047986
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                                                                                                                                                                             (UYCI-) UNIV CINCINNATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
les 36; Conserv
                                                                                                                                                                                                                                 Balasubramaniam A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36 AA;
                                                                         25-MAR-1998;
                                                                                                                            25-MAR-1998;
                        04-APR-2000,
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                                                                                                                                                                                                                                                                                                                                                                                                                       diarrhea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes the novel use of a composition containing a neuropeptide Y (RPY) Y4 receptor ligand for treatment of human reproductive disorders. The products described in the invention have endocrine, gynecological and cytostatic activity and can be used for the treatment of reduced reproductive function, delayed puberty, supranormal function of the reproductive organs, precoclous puberty, endometriosis, polycystic ovary syndrome, benign prostatic hyperplasia, impaired reproductive function or amenorheae. This sequence represents the human PPY neuropeptide which is used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PYY; peptide YY; human; electrolyte secretion; fluid secretion; nutrient uptake; lipolysis; vasoconstriction; gastrointestinal disorder; diarrhoea; Crohn's disease; irritable bowel syndrome; ileostomy;
                                                                         neuropeptide Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Compositions containing a neuropeptide Y Y4 receptor ligand selective for the hyporthalamic-plunitary-gonadal axis, for trearment of reproductive disorders e.g. delayed or precocious puberty, endometriosis and benign prostatic hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                   Neuropeptide; human; treatment; reproductive disorder; neurope endocrine; gynecological; cytostatic; puberty; endometriosis; polycystic ovary syndrome; prostatic hyperplasia; amenorrheae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 194; DB 21; 100.0%; Pred. No. 2.7e-20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Akinsanya K, Hayward A, Broqua P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 2; 12pp; English.
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                     Human neuropeptide PPY fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY87550 standard; peptide; 36
                                                                                                                                                                                                                                                                                                                                     98GB-0025969
                                                                                                                                                                                                                                                                                                                                                                                         98GB-0025969
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human peptide YY (PYY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-331548/29
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Matches 36; Conserv
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                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                         26-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                     26-NOV-1998;
                                                                                                                                                                                                                                 GB2344050-A.
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                                                                                                                                                                                                                                                                                  31-MAY-2000
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36 AA;
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         16-AUG-2001
                                                                                                                                                                                                                                                                                                                                              36;
                                                                                         Croom WJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU85987;
                                                                                                                                                                                                                                                                                                       Seguence
                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 10
AAU85987
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                                                                                                                                                                                                                                                                  The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/Nydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treament of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases in the action of peptidases to increase length of activity (half interference with physiological processes. ABB90823 to AAB92441 represent invention.
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                                                                                                                                                                                                         Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 36;
                                                                                                                                                                 Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 194; DB 22;
100.0%; Pred. No. 2.7e-20;
ive 0; Mismatches 0;
                                                                                                                                                                 Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                Disclosure; Page 327-328; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human peptide tyrosine-tyrosine (PYY).
                                                                                                                                                                 Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU06188 standard; peptide; 36 AA
                                                                                2000WO-US13576.
                                                                                                   99US-0134405
                                                                                                             99US-0153406
99US-0159783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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Best Local Similarity 100.
Matches 36; Conservative
                                                                                                                                            (CONJ-) CONJUCHEM INC
                                                                                                                                                                 Ezrin AM,
                                                                                                                                                                                   WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            36 AA;
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                                      WO200069900-A2
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          sapiens.
                                                                              17-MAY-2000;
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                                                                                                             10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2001
                                                                                                    17-MAY-1999;
                                                           23-NOV-2000
                                                                                                                                                                 Bridon DP,
                   Synthetic
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in the central nervous system (CNS). The method comprises administration of a peptide tyrosine-tyrosine (PYY) receptor agonist or a pancreatic polypeptide (PP) receptor agonist. The method is useful for the treatment of Alzheimer's disease and for reducing aluminium levels in the central nervous system, especially the brain, of a subject. The treatments are effective and do not impart excessive toxicological effects. The present sequence represents human PYY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hyperproliferative skin disease; hormone-dependent disease; homeostasis; intestinal disease; interleukin-8 production; smooth muscle contraction; feeding; blood pressure; pancreatic secretion; mutant; mutein; human; peptide YY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Increased biological potency, prolonged activity, increased half-life, glucose intolerance; insulin resistance; type II diabetes; bone disease; cancer; inflammatory disorder; obesity; developmental disorder;
                                                                                                                                                                                                                                                                                                                                                                                   treatment of Alzheimer's disease comprises administration of a peptide tyrosine receptor agonist or a pancreatic polypeptide receptor agonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method of reducing aluminium levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                     Reducing aluminium levels in the central nervous system, for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 7; 52pp; English.
                                                                                                                                      (UYNC-) UNIV NORTH CAROLINA STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU85987 standard; peptide; 36 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ő
                                                                                                                                                                                                           Taylor IL;
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07-FEB-2001; 2001WO-US03952
                                                                 08-FEB-2000; 2000US-0499980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-2002 (first entry)
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                                                                                                                                                                                                           Berg BM,
                                                                                                                                                                                                                                                                             WPI; 2001-550001/61
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The present sequence represents a human peptide yY (PYY). PYY triggers gain of function in glucose non-responsive foetal and adult islets which leads to glucose responsivity. The specification describes a method for promoting the growth of pancreatic cells and reducing degeneration of pancreatic tissue. The method comprises contacting pancreatic cells or tissue with a composition including PYY or an agonist of PYY. The method is used for treating a disease, especially in a human, associated with altered glucose metabolism, especially insulin resistance, glucose intolerance or glucose non-responsiveness, hyperglycemia, obbesity, hyperlipidemia, hyperfiltration or type II diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                            Promoting the growth of pancreatic cells and reducing degeneration of pancreatic tissue for treating a disease associated with altered glucose metabolism comprises contacting with a composition including (an agonist of) peptide YY .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; colon cancer; colon cancer antigen; diagnosis; detection; colorectal carcinoma; chromosome 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human colon cancer antigen protein SEQ ID NO:6128.
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100.0%; Pred. No. 8.9e-20;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                             Disclosure; Page 82-83; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0157137.
99US-0163280.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ruben SM, Barash SC,
                       (ONTO-) ONTOGENY INC.
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                                                                                                             WPI; 2000-565257/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97 AA;
                                                                                                                                       N-PSDB; AAA59713
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                                                                   Lu H;
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03-NOV-1999;
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                                                                 Pang K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 12
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    δλ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to modified biological peptides with increased biological potency, prolonged activity and/or increased half-life. The peptides of the invention are useful in the treatment of glucose intolerance which may be associated with inculin resistance pathologies, and in the treatment of type II diabetes. They are also useful for treating bone diseases, cancer, diseases related to inflammatory responses, obesity, autism, pervasive developmental classers, hyperproliferative skin diseases, incubition of They can be used for regulating blood glucose, enhancing mucosal regeneration in patients with intestinal diseases, inhibition of interleukin-8 production, stimulation of acid release, homeostasis, regulation of exocrine and endocrine secretions, smooth muscle contraction, feeding, blood pressure, body temperature and cell growth, regulation of food intake and energy balance, and stimulation of panceatic secretion or cell growth. AAU86019 represent the modified biological peptides of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide yY: PYY; pancreatic cell growth; pancreatic tissue degeneration; glucose metabolism; insulin resistance; glucose intolerance; glucose non-responsiveness; hyperglycemia; obesity; hyperlipidemia; hyperfiltration; type II diabetes mellitus.
                                                                                                                                                                            Novel modified biological peptide with increased biological potency, prolonged activity, increased half-life, for treating glucose intolerance associated or not with insulin resistance pathologies, type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 194; DB 23;
Pred. No. 2.7e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acid sequence of a human peptide yY (PYY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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29..97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "mature protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB08020 standard; Protein; 97 AA
                                                                                                                                                                                                                                                                                          Claim 5; Page 62; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
100.0%;
                                            (THER-) THERATECHNOLOGIES INC
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02-AUG-2000; 2000US-222619P
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Best Local Similarity 100.
Matches 36; Conservative
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                                                                                                                                     WPI; 2002-206179/26
                                                                                        Habi A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        36 AA;
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                                                                                                                                                                                                                                                     diabetes
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Sequence

AAB08020;

RESULT 11

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Peptide

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Gaps

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Length 97; Indels /

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           cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients of P by expressing by inserting the nucleic acids into a host cell and culturing the cell cexpressing the proteins. N and P peace when the prevention, diagnosis and treatment of colorectal carcinoms and cancers. Ashly196 to Ashl37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                 present invention.

N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present for SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon
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                                                                                                                                                                                                                                                                                                                                                                                             Length 176;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Parathyroid hormone (PTH) related peptide SEQ ID NO:283.
                                                                                                                                                                                                                                                                                                                                                                                             Score 194; DB 22;
Pred. No. 1.8e-19;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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100.0%;
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99US-0153406.
99US-0159783.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-JUN-2001 (first entry)
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Best Local Similarity 100.
Matches 36; Conservative
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                                                                                                                                                                                                                                                                                                                                                           176 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200069900-A2
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10-SEP-1999;
15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                             Sequence
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bonds with amino/Nydroxyl/thiol groups on blood components to form a benefidase stabilised therapeutic peptide composed of 3-50 amino acids. It are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptidas to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases. ABB90829 to AAB92441 represent invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (II) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity
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                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thibaudeau K;
                                                                                                                                                                                                                                                                                                      Length 36;
                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                   Score 188; DB 22;
Pred. No. 1.9e-19;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                        1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRPRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 329; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB91226 standard; Peptide; 36
                                                                                                                                                                                                                                                                                                    96.9%;
97.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0134406.
99US-0153406.
99US-0159783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-MAY-2000; 2000WO-US13576.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                     Local Similarity 97.2 es 35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide YY SEQ ID NO:400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bridon DP, Ezrin AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                    36 AA;
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Synthetic.
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15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB91226;
                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 14
AAB91226
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      Endogenous therapeutic
in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases intracellular uptake and interference with physiological processes. AAB90829 to AAB92441 represent peptides which can be used in the exemplification of the present invention.
                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gastro-enterological disorders. As part of a therapeutic composition they could be used for decreasing excess intestinal and augmenting nutrient transport, and for regulating cell proliferation and blood flow.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide YY; PYY; gastro-enterological disorders; intestinal water; electrolyte secretion; cell proliferation; nutrient transport; lipolysis; blood flow regulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR62049 describes the amino acid sequence of porcine peptide YY (PYY), which was isolated from the endocrine cells of the porcine gastrointestinal tract and panoreas. Using this sequence as a base the PYY analogues described in AAR62051-R62082 were produced. The new peptides were found to have a variety of properties, that made them useful as therapeutic agents in the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide derivs. - useful as therapeutic agents, for treating gastro-enterological disorders
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                                                                                                                                                                                                      Score 185; DB 22; Length 36;
Pred. No. 5e-19;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                       AAR62049 standard; peptide; 36 AA.
                                                                                                                                                                                                      95.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-JUN-1995 (first entry)
                                                                                                                                                                                                                  Best Local Similarity 94.4
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Porcine peptide YY (PYY).
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                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                        RESULT 15
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Score 184; DB 15; Length 36; Pred. No. 7e-19;

94.8%; 94.4%;

Query Match Best Local Similarity

Search completed: February 27, 2003, 14:36:15 Job time : 36 secs

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1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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ilarity 100.0%; Pred. No. 8.7
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/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd
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1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY
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US-09-229-900-2
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US-07-882-923-3
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US-09-181-941-1
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Listing first 45 summaries
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                                                                                                                OM protein
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Sequence 3, Al
Sequence 7, Al
Sequence 10, Sequence 13, Al
Sequence 1, Al
Sequence 6, M
Sequence 6, M
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Sequence 6, M
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Sequence 11,
  Sequence Sequence Sequence Sequence Sequence
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Sequence 2, Application US/08338395

Sequence 2, Application US/08338395

Patent No. 5574010

GENERAL INFORMATION:
APPLICANT: McFadden, David W

TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS WITH
TITLE OF INVENTION: PEPTIDE YY AND ANALOGS THEREOF
NUMBER OF SEQUENCES: 5

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Version #1.25
         US-08-994-946A-6
US-09-221-940-1
US-09-221-941-3
US-09-181-941-3
US-09-125-138-10
US-09-125-138-10
US-07-76-272-23
US-08-08-221-18-10
US-07-76-272-18
US-07-16-272-18
US-07-18-10-3
US-08-264-030-3
US-08-264-030-3
US-08-264-030-10
                                                                                                                                                                                                                                                                                                                                                  AUDRESSEE: POMS, SMITH, LANDE & ROSE STREET: 2029 Century Park East 38th Floor CITY: Los Angeles STATE: CA CONWROW.
                                                                                                                                                                                                                           ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Pacentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/338,395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Oldenkamp, David J
REGISTRATION NUMBER: 29421
REFERENCE/DOCKET NUMBER: 1070
TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-788-5446
TELEFAX: 310-277-1297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: HUMAN PEPTIDE YY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 310-277-1297
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acids
TYPE: amino acid
 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
TOPOLOGY: lin
MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                   USA
 144
144
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Gaps

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Length 36;

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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MB storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/054,393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 194; DB 3;
100.0%; Pred. No. 8.7e-22;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Balasubramanian, Ambikaipakan
TITLE OF INVENTION: PEPTIDE YY ANALOGS
CORRESPONDENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Frost & Jacobs, L.L.P.
ADDRESSEE: Z500 PNC Center, 201 East Fifth St.
CITY: Cincinnati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/047,986B
FILING DATE: 25 March 1998
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Kristyne A. Bullock
REGISTRATTON NUMBER: 9183030/508
TELECOMMUNICATION INFORMATION:
TELEPHONE: (513) 651-6731
TELEPHONE: (513) 651-6731
TELEFAX: 21-4396 FAJ CIN
TELEFEX: 21-4396 FAJ CIN
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                               BMR350/48000
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-047-986B-2
Sequence 2, Application US/09047986B
Patent No. 6046167
GENERAL INFORMATION:
                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION UNDRER: 36,912
REFERENCE/DOCKET 100HBER: BMR35
TELECOMMUNICATION INFORMATION:
TELEFAX: (202)639-6585
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: not relevant TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0
Best Local Similarity 100.0
Matches 36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 36 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 36 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                           CLASSIFICATION:
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                                                                                                                                                       FILING DATE:
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Patent No. 6017879
GENERAL INFORMATION:
APPLICANT: Lacroix, Jean S.
APPLICANT: Grouzmann, Eric
TITLE OF INVENTION: Template Associated NPY Y2-Receptor
TITLE OF INVENTION: Agonists
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESSE: Vanson & Elkins LLP
STREET: 1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 36;
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100.0%; Pred. No. 8.7e-22;
Live 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PS/2 Model 502 or 55SX OPERATING SYSTEM: MS-DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/329,151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Faul T. Clark
RECISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00537/105001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/038,534
FILING DATE: 3/29/93
APPLICATION NUMBER: 08/109,326
FILING DATE: 08/19/93
ATTORNEY/AGENT INFORMATION:
                                                                             GENERAL INFORMATION:
APPLICANT: Balasubramaniam, A.
TITLE OF INVENTION: ANALOGS OF PE
TITLE OF INVENTION: THEREOF
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ANDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                        Sequence 2, Application US/08329151
Patent No. 5604203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (617) 542-5070
(617) 542-8906
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Best Local Similarity 100.0
Matches 36; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                           Boston
Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: N/A
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington
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                      US-08-329-151-2
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  RESULT 2
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         WITH PEPTIDE YY AND ANALOGS THEREOF
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/14303
FILING DATE: 03 November 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/882,923
FILING DATE: 19920512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Score 194; DB 5;
; Pred. No. 8.7e-22;
0; Mismatches 0;
                                                                 ROSE
38th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Fitch, Even, Tabin & Flannery STREET: 4250 Executive Square, Suite 510 CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Boublik, Jaroslav H.
APPLICANT: Rivier, Jean E.F.
APPLICANT: Brown, Marvin R.
APPLICANT: Scott, Neal A.
TITLE OF INVENTION: NPY PEPTIDE ANALOGS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/503,198
                                                             ADDRESSEE: POMS, SMITH, LANDE & STREET: 2029 Century Park East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                         FILING DATE: 03 NOVEMBER 1995 CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Oldenkamp, David J
REGISTRATION NUMBER: 29421
REFERENCE/DOCKET NUMBER: 1070
TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-277-1297
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/07882923; Patent No. 5328899; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUMAN PEPTIDE YY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 36 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                               CORRESPONDENCE ADDRESS
                                                                       STREET: 2022 CITY: LOS Angeles
                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                         ZIP: 90067
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                                                                                                                                           COUNTRY:
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                                                                                                                                                                                                                                                                               Sequence 2, Application US/09229900
Patent No. 6288029
GENERAL INFORMATION:
APPLICANT: Mutter, Manfred
APPLICANT: Lacroix, Jean S.
APPLICANT: Grouzmann, Eric
TITLE OF INVENTION: Template Associated NPY Y2-Receptor
TITLE OF INVENTION: Agonists
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Vinson & Elkins LLP
                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/229,900 FILING DATE:
                                                                               Score 194; DB 3;
Pred. No. 8.7e-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 194; DB 4;
Pred. No. 8.7e-22;
Mismatches 0;
                                                                                                                                                         1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                             1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                             100.0%; Score 194; D
100.0%; Pred. No. 8.7
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Vinson & Elkins LLP
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY, AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFRERACE/POCKET NUMBER: BMR350/48000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)639-6585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application PC/TUS9514303 GENERAL INFORMATION:
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100.0%; Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McFadden, David W
TITLE OF INVENTION: TREATMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SS: not relevant not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (202)639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: Flodov At
                                                                                                 Best_Local Similarity 100.
Matches 36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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Best Local Similarity
Matches 36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washir
STATE: D.C.
COUNTRY: U.S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS
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US-09-229-900-2
                                         US-09-047-986B-2
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                                                                               Query Match
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                                                                                                                                                             Sequence 1, Application US/08329151
Patent No. 5604203
GENERAL INFORMATION:
APPLICANT: Balasubramaniam, A.
TITLE OF INVENTION: THEREOF
VUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS: ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1 Application US/09047986B
Fatent No. 6046167
GENERAL INFORMATION:
APPLICANT: Balasubramanian, Ambikaipakan
TITLE OF INVENTION: PEPTIDE YY ANALOGS
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Frost & Jacobs, L.L.P.
STREET: 2500 PNC Center, 201 East Fifth St.
CITY: Cincinnati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 184; DB 1;
Pred. No. 2.6e-20;
L; Mismatches 1
                                            1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PS/2 Model 502 or 555X OPERATING SYSTEM: MS-DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1) CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/329,151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          00537/105001
                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/038,534
FLING DATE: 3/29/93
APPLICATION NUMBER: 08/109,326
FILING DATE: 08/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (617) 542-5070
(617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.88;
94.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                              STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
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Best Local Similarity
Matches 34; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
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STREET: 22.
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US-08-329-151-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: McFadden, David W
TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS WITH
TITLE OF INVENTION: PEPTIDE YY AND ANALOGS THEREOF
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: POMS, SMITH, LANDE 6 ROSE
                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                          Score 184; DB 1; L
Pred. No. 2.6e-20;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: POMS, SMITH, LANDE & ROSE
2029 Century Park East 38th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRICE APPLICATION DATA:
APPLICATION NUMBER: US 07/219,596
FILING DATE: 15-0UL-1988
ATTOMESTAGENT INFORMATION:
NAME: Schumean, James J.
REGISTRATION NUMBER: 20,856
REFERENCE/DOCKET NUMBER: 52864
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION 1800RATION:
TELECOMUNICATION 1800RATION:
TELECOMUNICATION:
TELECOMUNICATION 1800RATION:
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FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/08338395
Patent No. 5574010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Oldenkamp, David J
REGISTRATION NUMBER: 29421
REFERENCE/DOCKET NUMBER: 1070
TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-788-5046
TELEFAX: 310-277-1297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   porcine peptide YY
                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 3:
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Best Local Similarity 94.4%;
Matches 34; Conservative
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30-MAR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34; Conservative
                                                                                                                                                                                                                                                                            ; TYPE: ANINO ACID
; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
US-07-882-923-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 2029 CC.
CITY: Los Angeles
STATE: CA
COUNTRY: USA
ZIP: 90067
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Best Local Similarity
Matches 34; Conserv
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 0 CLASSIFICATION:
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TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                            COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MB storage COMPUTER: IBM compatible OPERATING SYSTEM: MS-DOS SOFTWARE: Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/047,986B
FILING DATE: 25 March 1998
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Kristyne A. Bullock
REGISTRATION NUMBER: 42,371
REFERENCE/DOCKET NUMBER: 9183030/508
TELECOMUNICATION INFORMATION:

MACHINE REFERENCE/DOCKET NUMBER: 9183030/508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIF: 10036-2011
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: 1BM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASTSEO for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94.8%; Score 184; DB 3; L
94.4%; Pred. No. 2.6e-20;
live 1; Mismatches 1;
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NAME: COTUZZI, Laura A
REGISTRATION UNMBER: 30,742
REFERENCE/DOCKET NUMBER: 3909-0021-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/09181941
Patent No. 6440690
GENERAL INFORMATION:
APPLICANT: Mor, Amram
Vouldoukis, loannis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 29-JUN-1995
                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (513) 651-6731
TELEFAX: (513) 651-6981
TELEX: 21-4396 F&J Cin
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 94.4
Matches 34; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
            COUNTRY: USA
ZIP: 45202-4182
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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Pred. No. 2.6e-20;
1; Mismatches 1; Indels
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Sequence 1, Application PC/TUS9514303

GENERAL INFORMATION:
APPLICANT:
APPLICANT:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
WITH PEPTIDE YY AND ANALOGS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PLIAND DATE: PCT/1595/14303
FILING DATE: 03 November 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94.8%; Score 184; DB 5;
94.4%; Pred. No. 2.6e-20;
tive 1; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: POMS, SMITH, LANDE & ROSE
STREET: 2029 Century Park East 38th Floor
                                                                                                                                                                                                                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                   1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
; MOLECULE TYPE: No. 6440690e
; SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-181-941-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Oldenkamp, David J
REGISTRATION NUMBER: 29421
REFERENCE/DOCKET NUMBER: 107012F
TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-788-5046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release 11
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 36 amino acid
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    porcine peptide YY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                               94.88;
94.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 310-277-1297
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                           Query Match
Best Local Similarity 94.44
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 36 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 94.4'
Matches 34; Conservative
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Gaps
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Patent No. 5612454
GENERAL INFORMATION:
APPLICANT: Reminuma, Toshihiko
APPLICANT: Tajima, Masahiro
TITLE OF INVENTION:
Process for Purification of Polypeptide
NUMBER OF SEQUENCE: 31
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        83.5%; Score 162; DB 4; Length 36; 77.8%; Pred. No. 4.3e-17; Live 5; Mismatches 3; Indels
                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/776,272
                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LGUTA A
REGIESTRATION NUMBER: 30,742
REFERRNCE/DOCKET NUMBER: 3309-0021-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mueller & Player
P.O. Box 18218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-JUN-1995
                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
CLASSIFICATION: <u >Unknown></u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: No. 6440690e SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Wegner, Cantor, Mueller STREET: 1233 20th St. N.W. P.O. Bo CITY: Washington STATE: District of Columbia COUNTRY: United States of America 21P: 20036-8218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PC-DOS/MS-DOS
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TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                 MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 1:
                                                                                                          COMPUTER READABLE FORM:
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                                                              COUNTRY: USA
ZIP: 10036-2811
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                    CITY: New York
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US-07-776-272-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28;
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TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                              PEPTIDE YY AND USES
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CITY: Boston
STATE: Massacchusetts
COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READALE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 502 or 555X
OPERATING SYSTEM: MS-DOS (Version 5.0)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TURNEL ACLES.
NAME: POLIT: Clark
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/329,151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/038,534
FILING DATE: 3/29/93
APPLICATION NUMBER: 08/109,326
FILING DATE: 08/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Mor, Amram
Vouldoukis, Ioannis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/09181941 Patent No. 6440690 GENERAL INFORMATION:
                                       Sequence 9, Application US/08329151
Patent No. 5604203
                                                                                                   APPLICANT: Balasubramaniam, A. TITLE OF INVENTION: ANALOGS OF TITLE OF INVENTION: THEREOF
                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (617) 542-5070 (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 542-890.
TELEX: 200154
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear FEATURE:
                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 14
US-09-181-941-1
                  US-08-329-151-9
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Search completed: February 27, 2003, 14:38:41 Job time : 15 secs

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Sequence 2, Appli
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Sequence 6, Appli
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Sequence 3, Appli
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Sequence 6, Appli
Sequence 16, Appli
Sequence 16, Appli
Sequence 27, Appli
Sequence 27, Appli
Sequence 27, Appli
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(without alignments)
113.165 Million cell updates/sec
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/cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
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5.1.3
Compugen Ltd.
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2 US-10-038-045-4

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2 US-10-038-045-2

2 US-10-038-045-5

2 US-10-038-045-5

2 US-10-002-048A-2

US-10-002-048A-2

US-10-016-969-4

2 US-10-016-969-1

2 US-10-016-969-1

2 US-10-016-969-1

2 US-10-038-045-6

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2 US-09-955-528-16

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GenCore version
Copyright (c) 1993 - 2003
                                                                                                                                                                                                                                                                                                                                                                                       Published_Applications_AA:*
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Maximum Match 100%
Listing first 45 summaries
                                                   OM protein - protein search, using sw model
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length: 2000000000
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Match Length DB
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1178
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65:
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110:
113:
114:
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Maximum DB
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GENERAL INFORMATION:
APPLICANT: Amylin Pharmaceuticals, Inc.
APPLICANT: Amylin Pharmaceuticals, Inc.
APPLICANT: Pittner, Richard
APPLICANT: Potentii, James
APPLICANT: Paterniti, James
ATTLE OF INVENTION: Deptide YY and Peptide YY Agonists for the Treatment of Metabo TITLE OF INVENTION: Disorders
FILE REFERENCE: 24001-010
CURRENT APPLICATION NUMBER: US/10/016,969
CURRENT FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/256,216
WINDER OF FILING DATE: 2000-12-15
                                          Sequence 101, App.
Sequence 11, App.
Sequence 12, App.
Sequence 65, App.
Sequence 66, App.
Sequence 67, App.
Sequence 11, App.
Sequence 17, App.
Sequence 57, App.
Sequence 57, App.
Sequence 7, App.
Sequence 65, App.
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                                                                                                                                                                                           Sequence 14, Appl
Sequence 20, Appl
Sequence 39, Appl
Sequence 74, Appl
                          Sequence 46, Appl
Sequence 46, Appl
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    Sequence
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) US-09-205-658-58

) US-09-844-3534-58

US-09-844-3534-66

US-09-844-3534-46

US-09-844-3534-46

US-09-844-3534-101

US-09-844-3534-101

US-09-840-243B-11

US-09-840-243B-12

US-09-832-129-65

US-09-832-129-65

US-09-832-129-67

US-09-832-129-67

US-09-832-129-67

US-09-872-129-67

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US-09-872-129-67

US-09-872-129-67

US-09-873-1114-2

US-09-873-57

US-09-978-711A-4

US-09-978-711A-2

US-09-980-602-65

US-09-808-602-65

US-09-808-602-65

US-09-810-769-14

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US-09-810-769-14

US-09-810-769-14
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100.0%; Pred. No. 2.4e-19;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.0
SEQ ID NO 2
LENGTH: 36
  Ouery Match
Best Local Similarity 100.
Matches 36; Conservative
  TYPE: PRT
ORGANISM: Homo sapiens
  US-10-016-969-2
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APPLICANT: Mor, Amram Vouldoukis, Loannis Vouldoukis, Pierre Uicolas, Pierre TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION

Sequence 1199, Ap

Sequence 4, Application US/10038045 Patent No. US20020150964A1 GENERAL INFORMATION:

RESULT 2 US-10-038-045-4

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Gaps
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Nicolas, Pierre
TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
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                                                                                                              Indels
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COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,10/038,045
FILING DATE: 02-Jan-2002
CLASSIFCATION: CURKNOWN
PRIOR APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-UNN-1995
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
                                                                   Score 180; DB 12;
Pred. No. 1.6e-17;
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Pred. No. 4.2e-15;
5; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 3909-0021-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFRX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                            Mismatches
                                                                                                                                                                             1 IKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 34
                                                                                                                                                       3 IKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
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MOLECULE TYPE: No. US20020150964Ale
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                      RESULT 4
US-10-038-045-1
; Sequence 1, Application US/10038045
; Patent No. US20020150964A1
; GENERAL INFORMATION:
APPLICANT: Mor, Amram
Vouldoukis, Ioannis
                                                               Query Match 92.8%; Scc
Best Local Similarity 100.0%; P.
Matches 34; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 36 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      83.5%;
77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 10036-2811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
    ; ORGANISM: Homo sapiens
US-10-016-969-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 28; Conserv
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Patent No. US20020141985A1
GREERAL INFORMATION:
APPLICANT: Anylin Pharmaceuticals, Inc.
APPLICANT: Pittner, Richard
APPLICANT: Pittner, Richard
APPLICANT: Potentii, James
TITLE OF INVENTION: Peptide YY and Peptide YY Agonists for the Treatment of Metabolic
TITLE OF INVENTION: Disorders
FILE REFERENCE: 24001-010
CURRENT PELING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US/10/016,969
CURRENT FILING DATE: 2001-12-14
PRIOR FILING DATE: 2000-12-15
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OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
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                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/038,045
FILING DATE: 02-3an-2002
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.8%; Score 184; DB 12;
94.4%; Pred. No. 5.1e-18;
tive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 3909-0021-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-5556
TELEFAX: 650-493-5556
                                                            ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: NO. US20020150964Ale
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 36 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3
LENGTH: 34
TYPE: PRT
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                    NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
ULE TYPE: No. US:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 94.4
Matches 34; Conservative
                                                                                                            CITY: New York
                                                                                                                                                       USA
                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE
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GENERAL INFORMATION:
APPLICANT: Amylin Pharmaceuticals, Inc.
APPLICANT: Pittner, Richard
APPLICANT: Pittner, Richard
APPLICANT: Pittner, Richard
APPLICANT: Poung, Andrew
APPLICANT: Paterniti, James
TITLE OF INVENTION: Peptide YY and Peptide YY Agonists for the Treatment of Metabo
TITLE OF INVENTION: Disorders
FILE REFERENCE: 24001-010
CURRENT APPLICATION NUMBER: US/10/016,969
CURRENT FILING DATE: 2001-12-14
PRIOR FILLING DATE: 2000-12-15
                                                                                                                                                                         CORRUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSED for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/038,045
FILING DATE: 02-Jan-2002
CLASSIFICATION SATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 19-DEC-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-JUN-1995
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 146; DB 12;
Pred. No. 5.5e-13;
6; Mismatches 5;
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                            STREET: 1155 Avenue of the Americas
    ADDRESSEE: Pennie & Edmonds LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: No. US20020150964Ale SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 4, Application US/10016969
; Patent No. US20020141985A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 36 amino acids
                                                                                                                                                              MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 75.3%;
Best Local Similarity 69.4%;
Matches 25; Conservative
                                                                                                            ZIP: 10036-2811
COMPUTER READABLE FORM:
                                                 CITY: New York
                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens
US-10-016-969-4
                                                                                              COUNTRY:
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LENGTH: 36
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TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
                                                                                                                                                                                PEPTIDES FOR THE ACTIVATION OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
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OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
FILING DATE: 02-Jan-2002
CLASSIFICATION: CURKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.9%; Score 153; DB 12; 75.0%; Pred. No. 6.5e-14; iive 4; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNET/AGENT INFORMATION:
NAME: COTUZZI, LAURA
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 3909-0021-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4936
TELEFAX: 650-493-556
                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-UUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: No. US20020150964Ale SEQUENCE DESCRIPTION: SEQ ID NO: 2:
                                                                                                                                                         Nicolas, Pierre
TITLE OF INVENTION: PEPTIDES FOR
                                                                                                                                 Vouldoukis, Ioannis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mor, Amram
Vouldoukis, Ioannis
                                     Sequence 2, Application US/10038045
Patent No. US20020150964Al
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/10038045
Patent No. US20020150964Al
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
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CORRESPONDENCE ADDRESS:
                                                                                                                 Amram
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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les 27; Conservative
                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                            APPLICANT: Mor,
                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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RESULT 5
US-10-038-045-2
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Best Local Si
Matches 27;
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Length 36; Indels

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RESULT 10
US-10-016-969-5
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US-10-016-969-1
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                                           Gaps
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TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
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     Length 36;
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                                         Indels
                                                                                                                                                                                                   Sequence 2, Application US/10002048A
Publication No. US20020182616A1
GREERAL INFORMATION:
APPLICANT: Wahlestedt, Claes
APPLICANT: Daily, Bo.
TITLE OF INTENTION: Single Nucleotide Polymorphisms
FILE REFERENCE: 10806-143
CURRENT APPLICATION NUMBER: US/10/002,048A
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: SE 0004035-2
PRIOR FILING DATE: 2001-11-03
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Windows Version 2.0b
Score 144; DB 12;
Pred. No. 1e-12;
7; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    74.2%; Score 144; DB 9; I 66.7%; Pred. No. 3.2e-12; iive 7; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                             1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-UUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:

APPLICATION NUMBER: US./10/038,045
FILING DATE: 02-Jan-2002
CLASSIFICATION: <u >CURRENT</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/10038045 Patent No. US20020150964Al GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
   74.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
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   Query Match
Best Local Similarity 66.7'
Matches 24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIP: 10036-2811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
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Matches 24; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF
                                                                                                                                                                                      US-10-002-048A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-002-048A-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 2
LENGTH: 97
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APPLICANT: Paterniti, James
TITLE OF INVENTION: Peptide YY and Peptide YY Agonists for the Treatment of Metabo
TITLE OF INVENTION: Disorders
FILE REPERENCE: 24001-010
CURRENT APPLICATION NUMBER: US/10/016,969
CURRENT FILING DATE: 2001-12-14
PRIOR FILING DATE: 2000-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Pittner, Richard
APPLICANT: Young, Andrew
APPLICANT: Paternit, James
TITLE OF INVENTION: Peptide YY and Peptide YY Agonists for the Treatment of Metabo
TITLE OF INVENTION: Disorders
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Pred. No. 3.6e-11;
                                                                                                                                                                                                                                                                                                                                                             Score 141; DB 12;
Pred. No. 2.5e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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CURRENT APPLICATION NUMBER: US/10/016,969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/10016969
Patent No. US2002011985A1
GENERAL INCRMATION:
APPLICANT: Amylin Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Amylin Pharmaceuticals, Inc.
APPLICANT: Pittner, Richard
APPLICANT: Young, Andrew
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 39,742
REFERENCE/DOCKET NUMBER: 39
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/10016969 Patent No. US20020141985A1
                                                                                    TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                LENGTH: 36 amino acids
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                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
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66.7%;
                                                                                                                           TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 6
SOFWARE: Patentin version 3.0
SEQ ID NO 5
LENGTH: 34
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Best Local Similarity 63.9%;
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Matches 22; Conserv
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APPLICANT: LU, Dyung Alna M.
APPLICANT: PATTERSON, Chandra
TILLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES
FILE REPERENCE: PF-0701 USA
CURRENT APPLICATION NUMBER: US/09/965,528
CURRENT FILING DATE: 2001-09-26
PRIOR FILING DATE: 1999-05-19
PRIOR FILING DATE: 1999-07-15
PRIOR FILING DATE: 1999-07-15
PRIOR FILING DATE: 1999-07-15
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-10-04
NUMBER: OF SEQ ID NOS: 55
SOFTWARE: PERL PROGRAM
        SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: COURZEL, LAURA
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 3909-0021-999
TELECOMUNICATION INFORMATION:
TELEPHONE: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 52.1%; Score 101; DB 12; Best Local Similarity 73.9%; Pred. No. 2.9e-07; Matches 17; Conservative 4; Mismatches 2;
                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/038,045
FILING DATE: 02-Jan-2002
CLASSIFICATION: CURKNOWN>
                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/181,941
FILING DATE: 28-Oct-1998
APPLICATION NUMBER: US 08/574,701
FILING DATE: 19-DEC-1995
APPLICATION NUMBER: FR 95 07831
FILING DATE: 29-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: No. US20020150964Ale
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-10-038-045-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS
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BURFORD, Neil
BANDMAN, Olga
BAUGHN, Mariah R.
AZIMZAI, Yalda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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YUE, Henry
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-965-528-16
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LENGTH: 178
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APPLICANT:
APPLICANT:
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APPLICANT:
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Nicolas, Pierre
TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS
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                                                                                                                                                                                                                                        Length 36;
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S4.6%; Score 106; DB 12;
Best Local Similarity 48.6%; Pred. No. 1.1e-07;
Matches 17; Conservative 11; Mismatches 7;
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STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                  2 PLEPVYPGDNATPEQMAQYAADLKRYINMLTRPRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 PIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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TITLE OF INVENTION, NOVEL COMPOUNDS
FILE REFERENCE: GP-30015-C1
CURRENT APPLICATION NUMBER: US/09/757,712
CURRENT APPLICATION NUMBER: EP 97307187.1
PRIOR FILING DATE: 1997-09-16
PRIOR FILING DATE: 1997-09-16
PRIOR FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FASSESQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11; Mismatches
CURRENT FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/256,216
PRIOR FILING DATE: 2000-12-15
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09757712 Patent No. US20010016339A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/10038045
Patent No. US20020150964A1
GENERAL INFORMATION:
APPLICANT: Mor, Amram
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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                                                                                                                                                  TYPE: PRT
CORGANISM: Homo sapiens
US-10-016-969-1
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US-09-757-712-2
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Best Local Similarity
Matches 16; Conserv
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                                                                                                        SEQ ID NO 1
LENGTH: 36
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LENGTH: 95
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Length 23; Indels

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1 LOCATION: (30)
2 OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-300-1040
                                                                                                                                  Gaps
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US-09-925-300-1040
; Sequence 1040, Application US/09925300
; Patent No. US20020151681A1
; GENERAL INFORMATION:
; APPLICANT: Craig Rosen,
; APPLICANT: Steve Ruben
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
; CURRENT APPLICATION NUMBER: US/09/925,300
CURRENT FILLING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05988
; PRIOR FILLING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 1890
; SOFTWARE: PatentIN Ver. 2.0
; SEQ ID NO 140
; LENGTH: 85
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20020187523A1 5037143CD1
US-09-965-528-16
                                                                                   Ouery Match 51.0%; Score 99; DB 9; Length 178; Best Local Similarity 47.1%; Pred. No. 5.8e-06; Matches 16; Conservative 11; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      45.9%; Score 89; DB 10; Length 85; 66.7%; Pred. No. 5.2e-05; tive 6; Mismatches 1; Indels
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Best Local Similarity 66.7
Matches 14; Conservative
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ORGANISM: Homo sapiens
FEATURE:
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Search completed: February 27, 2003, 14:39:00 Job time : 13 secs

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Copyright (c) 1993
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OM protein - protein search, using sw model

February 27, 2003, 14:34:48; Search time 45 Seconds (without alignments) 76.908 Million cell updates/sec Run on:

Title: Perfect score: Sequence:

US-09-634-363-2 194 1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283224 seqs, 96134422 residues Searched:

Total number of hits satisfying chosen parameters:

283224

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

pIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
1: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES	ID Description	131358 periode YY - human	peptide YY pr	peptide YY	95 peptide YY	peptide YY	A60416 peptide YY - dog	peptide YY pr	A55914 peptide YY precurs	tic	pancreatic	pancreatic	pancreatic	neuropeptid		A41979 neuropeptide Y pre	neuropeptide Y		B30485 neuropeptide Y -		neuropeptide Y	neuropeptide Y				neuropeptide Y	×	peptide YY - ri	· · · · · · · · · · · · · · · · · · ·
	DB	. 7						7											~										
	Query Match Length 1	36	90	90	97	36	36	86	97	36	36	36	36	36	37	46	36	36	36	26	86	36	36	36	46	96	86	93	36
æ	Query Match	100.0	100.0	100.0	100.0	94.8	94.8	94.8	96.6	80.4	0	80.4	$^{\circ}$	8		75.8		74.2	74.2	74.2	74.2	m	ď.		2	ä	0	9.69	909
	Score	194	194	194	194	184	184	184	168	156	156	156	155	152	147	147	146	144	144	144	144	143	141	141	141	140	136	135	133
	Result No.		7	3	4	'n	9	7	æ	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	96

peptide YG - Ameri	pancreatic hormone													
A26781 S16943	PCBO	PCPG	PCDG	C61132	C60071	PCSH	PCCT	PCHU	A28256	A61132	D61132	B60413	A28578	JQ0365
0.0	-	Н	П	7	7	,-	ᆏ	-	7	~	m	7	7	7
37	36	36	93	36	36	59	99	95	126	36	36	36	36	36
68.0	58.2	57.2	57.2	54.6	54.6	54.6	54.6	54.6	54.6	54.1	54.1	53.6	53.6	53.1
132	113	111	111	106	106	106	106	106	106	105	105	104	104	103
30	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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peptide YY - human
C;Species: Homo sapiens (man)
C;Species: JanMar-1990 #sequence_revision 31-Dec-1991 #text_change 17-Mar-1999
C;Accession: A31358; A6066
R;Tatemoto, K.; Nakano, I.; Makk, G.; Angwin, P.; Mann, M.; Schilling, J.; Go, V.L.W.
Biochem: Biophys. Res. Commun. 157, 713-717, 1988
A;Title: Isolation and primary structure of human peptide YY.
A;Reference number: A31358
A;Molecule type: protein
A;Residues: 1-36 c4AP>
A;Ebrilein, G.A.; Eyselein, V.E.; Schaeffer, M.; Layer, P.; Grandt, D.; Goebell, H.;
Peptides: 10, 797-803, 1989
A;Title: A new molecular form of PYY: structural characterization of human PYY(3-36)
A;Reference number: A60676; MUID:90068171; PMID:2587421
A;Accession: A60676
A;Molecule type: protein
A;Residues: 1-36 c4BE>
C;Superfamily: pancreatic hormone
C;Keywords: amidated carboxyl end; intestine; neuropeptide
C;Keywords: amidated carboxyl end (Tyr) #status experimental
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Gaps ö Length 36; Indels Query Match 100.0%; Score 194; DB 2; Best Local Similarity 100.0%; Pred. No. 4.9e-19; Matches 36; Conservative 0; Mismatches 0;

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1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36 ò

1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36 qq

RESULT 2

934569
934569
954569
Peptide YY precursor (clone L2) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 09-May-1997
C;Accession: S34569
R;Kohrl, K; Nata Ara K; Yasa H; Nagai, A.; Konno, K.; Okamoto, H.
Biochim: Biophys. Acta 1173, 345-349, 1993
A;Title: Cloning and structural determination of human peptide YY cDNA and gene. A;Reference number: S33795; MUID:93305732; PMID:8318545
A;Reference number: S33795; MUID:93305732; PMID:8318545
A;Accession: S34569
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-90 < KOM>
C;Superfamily: pancreatic hormone

Length 90; 100.0%; Score 194; DB 2; 100.0%; Pred. No. 1.4e-18; Query Match Best Local Similarity .. 0

Gaps

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peptide YY - dog
C:Species: Canis lupus familiaris (dog)
C:Species: Canis lupus familiaris (dog)
C:Species: Theb-1993 #sequence_revision 11-Feb-1993 #text_change 17-Mar-1999
C;Accession: A60416
R:Eysselein, V.E.; Eberlein, G.A.; Grandt, D.; Schaeffer, M.; Zehres, B.; Behn, U.; S Peptides 11, 111-116, 1990
A;Title: Structural characterization of canine PYY.
A;Reference number: A60416; MUID:90259843; PMID:2342986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Rattus norvegicus (Norway rat)
C;Date: 31-Dec-1988 *text_change 16-Jul-1999
C;Date: 31-Dec-1988 *text_change 16-Jul-1999
C;Date: 31-Dec-1988 *text_change 16-Jul-1999
C;Date: 31-Dec-1988 *text_change 16-Jul-1999
R;Arcasion: A37955; A29364; JT0416
R;Krasinski, S.D.; Wheeler, M.B.; Leiter, A.B.
Mol. Endocrinol: 5, 433-440, 1991
A;Title: Isolation, characterization, and developmental expression of the rat peptide A;Reference number: A37955; MUID:91367188; PMID:1890992
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A; Rosaidues: 1-98 < KRRA
A; Cross-references: GB: 557220; NID: 9235283; PIDN: AAB19752.1; PID: 9235284
R; Leiter, A.B.; Toder, A.; Wolfe, H.J.; Taylor, I.L.; Cooperman, S.; Mandel, G.; Goo
J. Biol. Chem. 262, 12984-12988, 1987
A; Title: Peptide YY. Structure of the precursor and expression in exocrine pancreas.
A; Reference number: A29364; MUID: 88007492; PMID: 3644598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Superfamily: pancreatic hormone
C;Keywords: amidated carboxyl end; hormone; intestine
F;36/Modified site: amidated carboxyl end (Tyr) #status experimental
   F;36/Modified site: amidated carboxyl end (Tyr) #status experimental
                                                                             Score 184; DB 1; Length 36; Pred. No. 1e-17;
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Pred. No. 1e-17;
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Pred. No. 3.2e-17;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                       1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                          1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
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A;Experimental source: colon
C;Superfamily: pancreatic hormone
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94.4%;
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ilarity 94.4%;
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Best Local Similarity 94.4%
Matches 34; Conservative
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Best Local Similarity 94.4%
Matches 34; Conservative
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Matches 34; Conserv
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A; Residues: 1-98 <LEI>
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R; Tatemoto, K.
Proc. Natl. Acad. Sci. U.S.A. 79, 2514-2518, 1982
A; Title: Isolation and characterization of peptide YY (PYY), a candidate gut hormone that A; Reference number: A01574; WUID:82222168; PMID:6953409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S33795
S3795
peptide YY (clone S) - human
C; Species: Homo sapians (man)
C; Date: 19-Nar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Jun-2000
C; Accession: S33795
R; Kohri, K:; Nata, K:; Yonekura, H:; Nagai, A.; Konno, K.; Okamoto, H.
Biochim, Biophys. Acta 1173, 345-349, 1993
A;Title: Cloning and structural determination of human peptide YY cDNA and gene.
A;Reference number: S33795; MUID:93305732; PMID:8318545
A;Accession: S33795
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-97 <KOH>
A;Residues: 1-97 <KOH>
A;Residues: GB:D13897; NID:9391723; PIDN:BAA02997.1; PID:9391724
C;Superfamily: pancreatic hormone
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Cibate: 02-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 09-May-1997
CiAccession: S34568
CiAccession: S34568
A;Title: Cloning and structural determination of human peptide YY cDNA and ciAccession: S34568
A;Reference number: S33795; MUID:93305732; PMID:8318545
A;Accession: S34568
A;Accessi
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 30-Apr-1982 #sequence_revision 30-Apr-1982 #text_change 23-Aug-1996
C;Accession: A01574
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           Gaps
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       0;
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Pred. No. 1.5e-18;
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                                                                             1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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Best Local Similarity 100.0%; Pred. No. 1.5
Matches 36; Conservative 0; Mismatches
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A;Residues: 1-36 CTAT>
C;Superfamily: pancreatic hormone
C;Reywords: amidated carboxyl end; hormone
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0
       Conservative
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36;
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80.4%;
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1 Similarity 75.08;
27; Conservative
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Best Local Similarity 75.0
Matches 27; Conservative
A; Reference number: A60022
                                            A; Molecule type: protein A; Residues: 1-36 <PAN1>
                                                                                                                                                                                                                                                A; Molecule type: protein A; Residues: 1-36 <PAN2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: protein A; Residues: 1-36 <KIM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                        A; Accession: A56893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pancreatic peptide Y - alligator gar
N'Alternate names: pancreatic hormone Y
C'Species: Lepisosteus spatula (alligator gar)
C'Species: Lepisosteus spatula (alligator gar)
C'Date: 30-Jun-1991 #sequence_revision 30-Jun-1991 #text_change 06-Dec-1996
C'Accession: 507215
R'Pollock, H.G.; Kimmel, J.R.; Hamilton, J.W.; Rouse, J.B.; Ebner, K.E.; Lance, V.; Rawi
Gen, Comp. Endocrinol. 67, 375-382, 1987
A'Reference number: 507215; MUD:88030594; PMID:3311873
A'Reference number: 507215; MUD:88030594; PMID:3311873
A'Recession: 507215
A'Residues: 1-36 <POL>
A'Residues: 1-36 <POL>
C'Superfamily: pancreatic hormone
C'Superfamily: pancreatic hormone; pancreas
C'Keywords: amidated carboxyl end; hormone; pancreas
F;1-36/Product: pancreatic peptide Y *status experimental
                                                                                                                                                                                                                                                     C; Accession: A55914
R; Herzog, H.; Hort, Y.; Schneider, R.; Shine, J.
R; Herzog, H.; Hort, Y.; Schneider, R.; Shine, J.
R; Herzog, H.; Hort, Y.; Schneider, R.; Shine, J.
R; Herzog, H.; Hort, Y.; Schneider, R.; Shine, J.
A; Title: Seminal plasmin: recent evolution of another member of the neuropeptide Y gene A; Reference number: A55914
A; Reference number: A55914
A; Residues: preliminary
A; Molecule type: DNA
A; Residues: 1-97 <HER>A; Residues: 1-97 <HER>A; Cross-references: GB:L37369; NID:g565219; PIDN:AAC37326.1; PID:g567100
C; Genetics:
A; Introns: 63/2; 90/2
C; Superfamily: pancreatic hormone
C; Keywords: neuropeptide
C; Keywords: neuropeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pancreatic peptide Y - spiny dogfish
N.Alternate names: pancreatic hormone Y
C.Species: Squalus acanthias (spiny dogfish)
C.Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 16-Jun-2000
C.Accession: A60022; A56893
R.Pan, J.2.; Shaw, C.; Halton, D.W.; Thim, L.; Johnston, C.F.; Fairweather, I.; Buchanan Regul. Pept. 35, 252, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A, Title: Isolation and primary structure of peptide Y from the pancreas of the spiny dog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                         Species: Bos primigenius taurus (cattle)
Date: 23-Mar-1995 #sequence_revision 05-Apr-1995 #text_change 16-Jul-1999
Accession: A55914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             86.6%; Score 168; DB 2;
llarity 83.3%; Pred. No. 4.1e-15;
Conservative 3; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 156; DB 1;
Pred. No. 5.2e-14;
5; Mismatches 4
                                         1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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               1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 80.4%;
1 Similarity 75.0%;
27; Conservative
                                                                                                                                                                                      peptide YY precursor - bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 30; Conserv
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Matches
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Figorion, J.M.; Balasubramaniam, A.; Hazon, N.
Endocrinology 128, 2273-2279, 1991
A; Fitle: Structural characterization and biological activity of a neuropeptide Y-rela A; Feference number: A49743; MUID:91209266; PMID:2019251
A; Accession: A49743
A; Molecule type: protein
A; Residues: 1-36 CON>
C; Superfamily: pancreatic hormone
C; Superfamily: pancreatic hormone
C; Keywords: amidated carboxyl end; hormone
E; 1-36/Product: pancreatic peptide Y #status experimental <MAT>
F; 36/Modified site: amidated carboxyl end (Tyr) #status experimental
                                                                                                          A;Title: The primary structure of peptide Y (PY) of the spiny dogfish, Squalus acanth A;Reference number: A56893; MUID:92405520; PMID:1526112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession: A26377
R; Kimmel, J.R.; Plisetskaya, E.M.; Pollock, H.G.; Hamilton, J.W.; Rouse, J.B.; Ebner, Biochem. Biophys. Res. Commun. 141, 1084-1091, 1986
A; Title: Structure of a peptide from coho salmon endocrine pancreas with homology to A; Reterence number: A26377; MUID:87128023; PMID:3545195
A; Accession: A26377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pancreatic peptide Y - smaller spotted catshark
N'Alternate names: neuropeptide Y-related peptide
C'Species: Scyliorhinus canicula (smaller spotted catshark, smaller spotted dogfish)
C'Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 07-May-1999
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A; Note: this reference is an abstract
R; Pan, J.Z.; Shaw, C.; Halton, D.W.; Thim, L.; Johnston, C.F.; Buchanan, K.D.
Comp. Biochem. Physiol. B 102, 1-5, 1992
A; Title: The primary structure of peptide Y (PY) of the spiny dogfish, Squalu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pancreatic peptide Y - coho salmon
C;Species: Oncorhynchus Kisutch (coho salmon)
C;Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 06-Dec-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                     A) Experimental Source: pancreas
A) Note: sequence extracted from NCBI backbone (NCBIP:114876)
C) Superfamily: pancreatic hormone
C; Neywords: amidated carboxyl end; hormone; pancreas
F;1-36/Product: pancreatic peptide Y **status experimental <MAT>
F;36/Modified site: amidated carboxyl end (Tyr) **status experimental
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Pred. No. 5.2e-14;
5; Mismatches 4
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Pred. No. 5.2e-14;
5; Mismatches 4;
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Query Match
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R;Jensen, J.; Conlon, J.M.
A;Title: Characterization of peptides related to neuropeptide tyrosine and peptide tyros
A;Reference number: S27054; MUID:93092973; PMID:1459125
A;Accession: S27054
A;Accession: S27054
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R;Blomqvist, A.G.; Soderberg, C.; Lundell, I.; Milner, R.J.; Larhammar, D.
Proc. Natl. Acad. Sci. U.S.A. 89, 2350-2354, 1992
A;Title: Strong evolutionary conservation of neuropeptide Y: sequences of chicken, goldf A;Reference number: A41979; MUID:92196116; PMID:1549597
A;Accession: A41979
A;Accession: A41979
A;Accession: A41979
A;Anolecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Accession: $26954
R; Conlon, J.M.; O'Harte, F.
F; FEBS Lett. 313, 225-238, 1992
A; Title: The primary structure of a PYY-related peptide from chicken intestine suggests
A; Reference number: $26954; MUID: 93076900; PMID: 1446739
A; Reference number: protein
A; Recassion: $26954
A; Accession: $26954
A; Accession: $26954
A; Accession: $26954
A; Residues: 1-37 < CON>
C; Superfamily: pancreatic hormone
C; Keywords: amidated carboxyl end; hormone; intestine; neuropeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide YY-related protein, intestinal - chicken N.Alternate names: neuropeptide Y homolog; peptide tyrosine-tyrosine-related protein
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                                                                                                                                                                                                                                                                                                                                   C;Species: Gadus morhua (Atlantic cod)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 31-Oct-1997
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C.Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C.Species: Gallus gallus (chicken)
C.Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 07-Feb-1997
                                                                 Gaps
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Score 155; DB 2; Length 36;
Pred. No. 7e-14;
4; Mismatches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       75.8%; Score 147; DB 2; 69.4%; Pred. No. 8.3e-13;
                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8; Mismatches
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  79.98;
                                                      27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 69.4 Matches 25; Conservative
                                                                                                                                                                                                                                                                                                            neuropeptide Y - Atlantic cod
  Query Match
Best Local Similarity
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                                                            Matches
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A:Residues: 1-97 <BLO>
A.Cross-references: GB:M87294; NID:g212458; PIDN:AAA48991.1; PID:g212459
A.Cross-references: GB:M87294; NID:g212458; PIDN:AAA48991.1; PID:g212459
A.Rote: sequence extracted from NCBI backbone (NCBIP:88404)
C;Function: neuropeptide inducing a number of behavioral effects including stimula A;Pescription: neuropeptide inducing a number of behavioral effects including stimula C;Superfamily: pancreatic hormone
C;Keywords: amidated carboxyl end; appetite; hormone; neuropeptide
C;Keywords: amidated carboxyl end; appetite; hormone; neuropeptide
F;1-29-64/Product: neuropeptide f*status predicted <AMI>F;29-64/Product: neuropeptide f*status predicted <AMI>F;65-97/Domain: carboxyl-terminal propeptide f*status predicted cTP>
F;65-97/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following
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Pred. No. 2.5e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75.8%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24; Conservative
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Search completed: February 27, 2003, 14:38:20 Job time: 46 secs

us-09-634-363-2.rsp

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5.1.3
Compugen Ltd.
GenCore version
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OM protein - protein search, using sw model

February 27, 2003, 14:16:23 ; Search time 29 Seconds Run on:

(without alignments) 51.488 Million cell updates/sec

Perfect score: Sequence:

US-09-634-363-2

1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 seqs, 41476328 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_40:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Description	hon	P01305 sus scrofa	Q9eps2 mus musculu	P10631 rattus norv		P80952 phyllomedus		P09473 lepisosteus	-	P29206 raja rhina	0918p2 brachydanio	Q9pta0 dicentrarch	_		P80167 gadus morhu		_		P28673 gallus gall		P09640 oryctolagus	P81028 oreochromis	homo	-	rattn	ovis	P29949 rana ridibu	P33689 xenopus lae		Q9pt98 dicentrarch	Q9pt99 dicentrarch	674 torpedo m	P48098 lampetra fl
SUMMARIES		ΠD	PYY_HUMAN	PYY_PIG	PYY_MOUSE	PYY_RAT	PYY_BOVIN	SPYY_PHYBI	PYY_AMICA	PYY_LEPSP	PYY_ONCKI	PYY_RAJRH	PYY_BRARE	NEUY_DICLA	NEUY_ONCMY	PYY_RANRI	NEUY_GADMO	NEUY_ICTPU	PYY_CHICK	NEUY_BRARE	NEUY_CHICK	NEUY_PIG .	NEUY_RABIT	PYY_ORENI	NEUY_HUMAN	NEUY_MOUSE	NEUY_RAT	NEUY_SHEEP	NEUY_RANRI	NEUY_XENLA	NEUY_CARAU	PY_DICLA		•	PYY_LAMFL
		ngth DB	97 1			98 1	. ,		36 1	36 1	36 1	36 1	97 1	99 1	36 1	36 1	36 1	95 1	37 1	96 1			36 1							97 1				98 1	93 1
ð	Query	Match Length	100.0						81.4			•	79.4	•	•			•	٠	٠	•		•	•	٠	•	•	ω.	'n.	ς.	ς.	ä	ö	70.1	6
		Score	194	184	184	184	168	162	158	156	155	155	154	154	153	153	152	149	147	147	147	146	144	144	144	144	144	143	141	141	140	139	137	136	135
	Result	No.	г	7	m	4	5	9	7	80	6	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33

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P09641 myoxocephal P48097 lampetra fl	P09475 lophius ame P80024 petromyzon	P01302 bos taurus P01300 sus scrofa	P01299 canis famil	P41337 larus argen	P41336 oryctolagus	
PYY_MYOSC NEUY_LAMFL	PPY_LOPAM PMY_PETMA	PAHO_BOVIN	PAHO_CANFA	PAHO_LARAR	PAHO_RABIT	PAHO_TAPPI
	~ ~				, ₍ ,	
36 104	69 36	131	93	30	36	36
68.6 68.6	68.0 62.9	58.2	57.2	55.2	54.6	54.6
133 133	132	113	111	107	106	106
34 35	36 37	38 36	40	4 5 4	44	45

ALIGNMENTS

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This SWISS-PROT entry is copyright. It is produced through a collaboration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=90068171; PubMed=2587421; MEDLINE=90068171; PubMed=2587421; Beridin G.A., Eysselein V.E., Schaeffer M., Layer P., Grandt D., Goebell H., Niebel W., Davis M., Lee T.D., Shively J.E., Reeve J.R. Jr.; Reeve J.R. Jr.; Raw molecular form of PYY: structural characterization of human PYY(1-36) and PYY(1-36).; Peptides 10:797-803(1989).

Peptides 10:797-803(1989).

Peptides 10:797-803(1989).

HAS A VASOCONSTRICTORY ACTION AND INHIBITIS JEJUNAL AND COLONIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -:- SUBCELLULAR LOCATION: Secreted.
-:- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A LONG FORM (SHOWN HERE) AND A SHORT FORM; ARE PRODUCED BY ALTERNATIVE SPLICING.
-:- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                                                                                                                                                        TISSUE-Colon mucosa;
MEDLINE-23305732, PubMed-8318545;
Kohri K., Nata K., Yonekura H., Nagai A., Konno K., Okamoto H.;
"Cloning and structural determination of human peptide YY cDNA and
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tatemoto K., Nakano I., Makk G., Angwin P., Mann M., Schilling J., Go V.L.W.:
                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (NOV-1993) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Isolation and primary structure of human peptide YY."; Biochem. Biophys. Res. Commun. 157:713-717(1988).
                                                    01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Peptide YY precursor (PYY) (Peptide tyrosine).
 97 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 29-64, AND SYNTHESIS OF 29-64. MEDLINE-89076307; PubMed=3202875;
                                                                                                                                                                                                                                                                                                               Biochim. Biophys. Acta 1173:345-349(1993).
 PRT;
                            01-MAR-1989 (Rel. 10, Created)
01-NOV-1995 (Rel. 32, Last seq
16-OCT-2001 (Rel. 40, Last anno
 STANDARD;
                                                                                                                                                                                                                                                                                                                                                             TISSUE=Lymphocytes;
                                                                                                                                                                                            [1]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
PYY_HUMAN
P10082;
                                                                                                                                                                                                                                                                                                                                                                                   Herzog H
                                                                                                                                                                                                                                                                                              gene.
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us-09-634-363-2.rsp

MIM;

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-:- SUBCELLULAR LOCATION: Secreted.
-:- SIMILARITY: BELONGS TO THE NPY / PPY / PXY FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOBILITY.
SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AF325866; AAG42908.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN=BALB/c; TISSUE=Liver;
                                                                                                                                                                            94.88;
                                                                                                                                                      36 AA; 4242 MW;
                                                                                                                                                                                                  34; Conservative
                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28
28
793
64
                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
                                                                                                                                Hormone; Amidation.
                                                                                                                                                                                      Local Similarity
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68
64
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                                                                                                                                                                                                                                                                                                  PYY_MOUSE
                                                                                                                                                      SEQUENCE
                                                                                                                                                                            Query Match
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SIGNAL
PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROPEP
MOD_RES
                                                                                                                                             MOD_RES
                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                           RESULT 3
PYY_MOUSE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Isolation and characterization of peptide YY (PYY), a candidate gut
                                                                                                                                                                                                                                                                 AMIDATION (G-65 PROVIDE AMIDE GROUP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sus scrofa (Pig), and Canis familiaris (Dog).

Canis familiaris (Dog).

Evaryora: Metazoca: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                        SMART; SM00309; PAH; 1.
PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
Hormone; Cleavage on pair of basic residues; Pancreas; Amidation;
                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                100.0%; Score 194; DB 1; Length 97; 100.0%; Pred. No. 1.1e-19;
                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                          MISSING (IN SHORT ISOFORM).
T -> R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hormone that inhibits pancreatic exocrine secretion."; Proc. Natl. Acad. Sci. U.S.A. 79:2514-2518(1982).
                                                                                                                                                                                                                                                                                                 /FTId=VAR_006382.
DD16B73407F656A4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Peptide YY (PYY) (Peptide tyrosine tyrosine).
                                                                                                                                                                                                                                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
                                                                                                                                                                                                                                                                                                                                                                                       29 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 64
                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. ....
                                                                                                                                                                                                                                           PEPTIDE YY.
an email to license@isb-sib.ch)
                                                                                                                                        Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
                                                                                                                               interPro; IPR001955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES=Pig;
MEDLINE=82222168; PubMed=6953409;
                  EMBL, D13897; BAA02997.1; --
EMBL, D13899; BAA03098.1; --
EMBL, D138992; BAA03000.1; --
EMBL, L25648; BAA36433.1; --
EMBL, A31358; A31358; PIR; A31358; A01358.
                                                                                                                                                                                                               Signal; Alternative splicing. SIGNAL
                                                                                                                                                                                                                                                                                                           11046 MW;
                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
nes 36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                        Genew; HGNC:9748; PYY.
                                                                                                                                                                                                                                          64
97
64
97
72
                                                                                                                                                                                                                                                                                                           97 AA;
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68
64
91
72
                                                                                                                     600781;
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                                                                                                                                                                                                                                                                MOD_RES
VARSPLIC
                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
send
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PYY_PIG
P01305;
                                                                                                                                                                                                                                           PEPTIDE
                                                                                                                                                                                                                                                                                       VARIANT
                                                                                                                                                                                                                                                      PROPEP
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Matches

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RESULT 2

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brown G.J., James R., Eddie L.W.;
Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
HAS A VASOCONSTRICTORY ACTION AND INHIBITIS JEJUNAL AND COLONIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AMIDATION (G-65 PROVIDE AMIDE GROUP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide YY precursor (PYY) (Peptide tyrosine tyrosine) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE: PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE: PS50276; PANCREATIC_HORMONE_2; 1.
HORMONE; Cleavage on pair of basic residues; Pancreas; Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 184; DB 1; Length 36; Pred. No. 7.7e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                          02CD6B8C586DCC8D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-0cT-2001 (Rel. 40, Created)
16-0cT-2001 (Rel. 40, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.4%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BY SIMILARITY. PEPTIDE YY.
                                                                                                                                                                                                                                                                         PROSITE; PS00265; PANCREATIC_HORMONE_1; 1. PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
PIR; A01574; YYPG.
PIR; A60416; A60416.
HSSP; P01303; 1RON.
InterPro; IPR001955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                              AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MGD; MGI:99924; Pyy.
InterPro; IPR001955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRINTS; PRO0278; PANCHORMONE.
Probom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
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1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
  Signal.
SIGNAL
PEPTIDE
                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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MOD_RES
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SIGNAL
                                                               PROPEP
                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                          RESULT 5
PYY_BOVIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=88321122; PubMed=3413293;
Corder R., Gaillard R.C., Boehlen P.;
"Isolation and sequence of rat peptide YY and neuropeptide Y.";
Regul. Pept. 21:253-261(1988).
- Pept. BUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
HAS A VASOCONSTRICTORY ACTION AND INHIBITIS JEJUNAL AND COLONIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDINE-91367188; PubMed-1890992; Krasinski S.D., Wheeler M.B., Leiter A.B.; Isolation, characterization, and developmental expression of the rat Mol. Endocrinol. 5:433-440(1991).
                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mandel G., Goodman R.H.; "Peptide YY. Structure of the precursor and expression in exocrine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
Hormone; Cleavage on pair of basic residues; Pancreas; Amidation;
                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE-88007492; PubMed-3654598;
Leiter A.B., Toder A., Wolfe H.J., Taylor I.L., Cooperman S.
Mandel G., Goodman R.H.;
                                                             Length 93;
                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                   01-JUL-1989 (Rel. 11, Created)
01-JUL-1989 (Rel. 11, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Peptide YY precursor (PYY) (Peptide tyrosine tyrosine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                      2320990F05992030 CRC64;
                                                           Score 184; DB 1;
Pred. No. 2.4e-18;
                                                                                                                                            1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                       98 AA
                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PIR; A29364; A29564.
PIR; A37955, A37955.
HSSP, P01303, IRON.
InterPrc; IPR001955; Pancreatic_hormp.
Pfam; PF00159; hormone3; 1.
PrinTs; PR00278; PANCTRORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               J. Biol. Chem. 262:12984-12988(1987).
                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: Secreted
                 10481 MW;
                                                           94.8%;
94.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; S57220; AAB19752.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA41222.1; -.
                                                                                                     34; Conservative
                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus (Rat)
                                                                                 Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 29-64.
93
93 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10116;
NON_TER
SEQUENCE
                                                               Query Match
                                                                                                                                                                                                                                                                                                PYY_RAT
P10631;
                                                                                                     Matches
                                                                                                                                                                                                                                                      RESULT 4
                                                                                                                                                                                                                                                                        PYY_RAT
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Proc. Natl. Acad. Sci. U.S.A. 92:594-598(1995).
-i- FUNCTION: THIS GUT PEPTIDE INHIBITS EXOCRINE PANCREATIC SECRETION,
HAS A VASOCONSTRICTORY ACTION AND INHIBITIS JEJUNAL AND COLONIC
                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoldea;
Bovidae; Bovinae; Bos
                                                                        AMIDATION (G-65 PROVIDE AMIDE GROUP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
HORMONE; Cleavage on pair of basic residues; Pancreas; Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AMIDATION (G-65 PROVIDE AMIDE GROUP).
B3A7B6A768BB3AE0 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIINE=95132646; PubMed=7831336;
Herzog H., Hort Y., Schneider R., Shine J.;
"Seminalplasmin: recent evolution of another member of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 168; DB 1; Length 97; Pred. No. 3.8e-16; 3; Mismatches 3; Indels
                                                                                                                                          Score 184; DB 1; Length 98
Pred. No. 2.6e-18;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-oCT-1996 (Rel. 34, Created)
01-oCT-1996 (Rel. 34, Last sequence update)
16-oCT-2001 (Rel. 40, Last annotation update)
Peptide YY precursor (PYY) (Peptide tyrosine tyrosine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -:- SUBCELLULAR LOCATION: Secreted.
-:- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                               994C0C3AD6A8A7DE CRC64;
                                                                                                                                                                                                                                         1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                     29 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 64
                                                                                                                                                                                                                                                                                                                                                                                                                  97 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BY SIMILARITY. PEPTIDE YY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Mismatches
                           PEPTIDE YY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
28
64
98
64
11121 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuropeptide Y gene family.";
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                                                                                                                                                 94.88;
                                                                                                                                                                      94.48;
                                                                                                                                                                                               34; Conservative
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97
64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   taurus (Bovine).
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30; Conserv
                                                                                                                                                                      Best Local Similarity
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68
68
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P51694;
                                                                      MOD_RES
SEQUENCE
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MEDLINE=91296574; PubMed=2067973;
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P09473;
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-:- SUBGELLULAR LOCATION: Secreted.
-:- TISSUE SPECIFICITY: SKIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptide family: isolation, structure, synthesis, and endocrine activity.";
                                                                                                                                                                                                                             Skin peptide tyrosine-tyrosine (SPYY).
Phyllomedusa blocolor (Two-colored leaf frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amia calva (Bowfin).
Usukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Amiiformes; Amiidae; Amia.
NCBI_TaxID=7924;
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                                                                                                                                                                                                                                                                                                                                                                                               TISSUE-Skin;
MEDLINE-95024100, PubMed-7937944;
Mor A., Chartrel N., Vaudry H., Nicolas P.;
"Skin peptide tyrosine-tyrosine, a member of the pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PSO0265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS0276; PANCREATIC_HORMONE_2; 1.
Amphibian skin; Fungicide; Hormone; Amidation. MOD_RES 36 36 MAIDATION (BY SIMILARITY).
SEQUENCE 36 AA; 4265 MW; 0725D316031827AE CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Proc. Natl. Acad. Sci. U.S.A. 91:10295-10299(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vouldoukis I., Shai Y., Nicolas P., Mor A.;
"Broad spectrum antibiotic activity of skin-PYY.";
FEBS Lett. 380:237-240(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           83.5%; Score 162; DB 1; 77.8%; Pred. No. 7.6e-16;
                                                                                                                                                               01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
29 YPAKPQAPGEHASPDELNRYYTSLRHYLNLVTRQRF 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001955; Pancreatic_hormn. Pfam; PF00159; hormone3; 1. PRINTS; PR00278; PANCHORMONE. ProDom; PD001267; Pancreatic_hormn; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-96186932; PubMed-8601432;
                                                                                                                                                                                                                                                                                                               Phyllomedusinae; Phyllomedusa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 77.8
hes 28; Conservative
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                                                                                                                             STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHARACTERIZATION.
                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=8393;
                                                                                                                      SPYY_PHYBI
P80952;
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SEQUENCE.
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                                                                                                     SPYY_PHYBI
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PYY_AMICA
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Gaps
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Fairweather I., Buchanan K.D.;

Fischation and primary structure of the peptide Y from the pancreas of the spiny dogfish, Squalus acanthias.";

Regul. Pept. 35:252-228(1991).

-i- FUNCTION: ELICITS AN INCREASE IN ARTERIAL BLOOD PRESSURE.

-i- FUNCTION: Secreted.

-i- SIBMELAULAR LOCATION: Secreted.

-i- SIBMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
Pollock H.G., Kimmel J.R., Hamilton J.W., Rouse J.B., Ebner K.E., Lance V., Rawitch A.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Isolation and structures of alligator gar (Lepisosteus spatula)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAR-1989 (Rel. 10, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Peptide YY-1ike (PYY) (Neuropeptide Y-related peptide).
Lepisosteus spatula (Alligator gar) (Atractosteus spatula),
Scyllorhinus canicula (Spotted dogfish) (Spotted catshark), and
Squalus acanthias (Spiny dogfish).
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"Structural characterization and biological activity of a
neuropeptide Y-related peptide from the dogfish, Scyliorhinus
canicula.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Actinopterygii; Neopterygii; Semionotiformes; Lepisosteidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 158; DB 1; Length 36;
Pred. No. 2.7e-15;
3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AMIDATION.
56B46F3C08666671 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36 AA.
                                                                                                                                                                                                                                                              PRINTS; PRO0278; PANCHORMONE.
ProDom: PD001267; Pancreatic_hormn; 1.
SMART; SM0309; PAH; 1.
PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      insulin and pancreatic polypeptide.";
Gen. Comp. Endocrinol. 67:375-382(1987);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE, AND SYNTHESIS.
SPECIES-S.canicula; TISSUE-Pancreas;
MEDLINE-91209266; PubMed=2019251;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPECIES=L.spatula; TISSUE=Pancreas;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81.4%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                   Hormone, Amidation.
MOD_RES 36
SEQUENCE 36 AA; 4333 MW;
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nes 28; Conservative
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Cyprinidae;
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P29206;
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Q9I8P2;
               Hormone; P
MOD_RES
SEQUENCE
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                                                                                                                                                                        PYY_RAJRH
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                                                                                                                                                                                                                                                                                                                            01-MAR-1989 (Rel. 10, Last sequence update)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-1001 (Rel. 40, Last annotation update)
16-OCT-1001 (PYY).
Oncorhynchus Mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi;
Actinopterygii: Neopterygii: Teleostei: Euteleostei:
Protacanthopterygii: Salmoniformes: Salmonidae: Oncorhynchus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kimmel J.R., Plisetskaya E.M., Pollock H.G., Hamilton J.W.,
Rouse J.B., Ebner K.E., Rawitch A.B.;
"Structure of a peptide from coho salmon endocrine pancreas with
homology to neuropeptide Y.";
Biochem. Biophys. Res. Commun. 141:1084-1091(1986).
                                                                                                                                                                                                ö
                                                                                                                                                                          Length 36;
                                                                                                                                                                                                4; Indels
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MEDLINE-93157164; PubMed-1494498;
Barton C.L., Shaw C., Halton D.W., Thim L.;
Rainbow trout (Oncorhynchus mykiss) neuropeptide Y.";
Peptides 13:1159-1163(1992).
-: SUBCELLULAR LOCATION: Secreted.
-: SUBLIARITY: BELONGS TO THE NPY / PPY FAMILY.
                                                                                                                                      AMIDATION.
56A6D8CC086660AA CRC64;
                                                                                                                                                                        DB 1;
                                                                                                                                                                       Score 156; DB 1;
Pred. No. 5e-15;
5; Mismatches
                                                                                                                                                                                                                                1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                    36 AA
                                                                                                  PROSITE; PS00265; PANCREATIC_HORMONE_1; 1. PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PIR: S07215; PCGXA.
PIR: A60022; PCDFY.
PIR: A49743; A49743.
HSSP: P01303; 1RON.
InterPro; IPR001955; Pancreatic_hormn.
Pfam: PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; PANCHORMONE.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIR; A26377; A26377.
HSSP; P01303; IRON.
InterPro; IPR001955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=0.kisutch; TISSUE=Pancreas;
MEDLINE=87128023; PubMed=3545195;
                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                        01-MAR-1989 (Rel. 10, Created)
                                                                                                                                                                       80.4%;
75.0%;
                                                                                                                                                 36 AA; 4291 MW;
                                                                                                                                                                      Query Match 80.4
Best Local Similarity 75.0
Matches 27; Conservative
                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                           Hormone; Amidation.
                                                                                                                                                                                                                                                                                                   PYY_ONCKI
P09474;
                                                                                                                                                 SEQUENCE
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                                                                                                             Gaps
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Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
                                                                                                                                                                                                                                                                                                                                                  Raja rhina (Skatė).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea; Hypnosqualea; Pristiorajea; Batoidea;
Rajiformes; Rajidae; Raja.
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                                                                           Length 36;
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                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                              AMIDATION.
56A6D8CC08666671 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AMIDATION.
07A7D9DC196660B6 CRC64;
                                                                           Score 155; DB 1; L
Pred. No. 6.9e-15;
4; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 155; DB 1;
Pred. No. 6.9e-15;
7; Mismatches 3;
                                                                                                                                                                                                                                                                                    01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Peptide YY precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                          97 AA.
                                                                                                                                                                                                                                                         36 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; PR001955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
PRINTS: PR00278; PANCHORMONE.
PRODOM; PD001267; PANCHORMONE.
PRODOM; PD001267; PANCHORMONE.
PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
                                                                                                                                                                                                                                                         PRT;
                                                                           79.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hormone, Amidation.
MOD_RES 36
SEQUENCE 36 AA; 4251 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             79.98;
72.28;
                              36 36
36 AA; 4305 MW;
                                                       Query Match
Best Local Similarity 75.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 72.2%
                                                                                                                                                                                                                                                         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                     Peptide YY-like (PYY).
               Amidation.
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SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=30478;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2: 1.
Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.
SIGNAL 1 28 BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jensen J., Conlon J.M.;

"Characterization of peptides related to neuropeptide tyrosine and peptide tyrosine-tyrosine from the brain and gastrointestinal tract of teleost fish.";

Eur. J. Biochem. 210:405-410(1992).

-!-FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN SECRETION OF GONADOTROPHIN-RELEASE HORMONE.

-!- SUBCELLULAR LOCATION: Secreted.

-!- SIMILARITY: BELONGS TO THE NPY / PPY / PXY FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AMIDATION (G-65 PROVIDE AMIDE GROUP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostel; Euteleostel;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID=8022;
      Cerda-Reverter J.M., Martinez-Rodriguez G., Zanuy S., Carrillo M.
                                                                                          (Dicentrarchus labrax).";
Submitted (APF-1998) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN
SECRETION OF GONADOTROPHIN-RELEASE HORMONE.
                                                          "Deduced peptide sequence of neuropeptide Y exon 2 from sea bass
                                                                                                                                                                                                         -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4EEFAED164964184 CRC64;
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Pred. No. 3.2e-14;
6; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Promis PRO0278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AJO05381; CAB64935.1; --
HSSP; PO1303: TRON
INTERPRO; IPRO01955; Pancreatic_hormn.
Pfam; PF00159; hormone3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HSSP; P01303; 1RON.
InterPro; IPR001955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AJ005378; CAB64932.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       79.48;
72.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99 AA; 11260 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29
68
64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Simil
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SEQUENCE
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      Db
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dicentrarchus labrax (European sea bass).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Percoidei;
                                                                                   "Zebrafish genes for neuropeptide Y and peptide YY reveal origin by chromosome duplication from an ancestral gene linked to the homeobox
            Soderberg C., Wraith A., Ringvall M., Yan Y.L., Postlethwait J.H., Brodin L., Larhammar D.; "Zebrafish genes for neuropeptide Y and peotide YV reveal research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cerda-Reverter J.M., Martinez-Rodriguez G., Zanuy S., Carrillo M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEPTIDE YY.

C-TERMINAL EXTENSION (BY SIMILARITY).

AMIDATION (G-65 PROVIDE AMIDE GROUP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
Hormone; Cleavage on pair of basic residues; Amidation; Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                          J. Neurochem. 75:908-918(2000).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (POTENTIAL).
96EA07EF0991AC2D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 154; DB 1;
Pred. No. 3.1e-14;
6; Mismatches 4,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NEUY_DICLA STANDARD; PRT; 99 AA. Q9PTAG; 09PT97; 16-OCT-2001 (Rel. 40, Created) 16-OCT-2001 (Rel. 40, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZFIN; ZDB-CENE-980526-71; pyy.
InterPro; IPR001955; Pancreatic_hormn.
Pfam; PR00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
PRODOM; PD001267; Pancreatic_hormn; 1.
SWART; SM00309; PAH; 1.
MEDLINE=20396163; PubMed=10936170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neuropeptide Y precursor (NPY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AF233875; AAF79942.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11175 MW;
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64
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nes 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P01303; 1RON
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29
68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE-Brain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide
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                                                                                                                                                     cluster."
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NEUY_DICLA

RESULT 12

Matches

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Gaps

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Length 99; Indels us-09-634-363-2.rsp

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1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                               Rana ridibunda (Laughing frog) (Marsh frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
NCBI_TaxID=8406;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gadus morhua (Atlantic cod).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conlon J.M., Chartrel N., Vaudry H.,
"Primary structure of frog PYY: implications for the molecular
evolution of the pancreatic polypeptide family.";
Peptides 13:145-149(1992).
-: SUBCELLULAR LOCATION: Secreted.
-: SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
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                                                                                                                                                                                   Indels
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                                                                                                         AMIDATION.
E2A32293A866611C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AMIDATION.
0725C3CC6304F663 CRC64;
                                                                                                                                               Score 153; DB 1; I
Pred. No. 1.3e-14;
5; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 78.9%; Score 153; DB 1; L
Best Local Similarity 75.0%; Pred. No. 1.3e-14;
Matches 27; Conservative 4; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-JUN-2002 (Rel. 41, Last annotation update)
Neuropeptide Y (NPY).
                                                                                                                                                                                                                                                                                                                                              01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-CCT-2001 (Rel. 40, Last annotation update)
Peptide YY-like (PYY).
Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORNONE.
ProDom; PD001267; Pancreatic_hormn; 1.
PR05ITE; PS00265; PANCREATIC_HORNONE_1; 1.
PR05ITE; PS50276; PANCREATIC_HORNONE_2; 1.
Mouropeptide, Amidation.
MOD_RES 36 36 36 AMIDATION.
SEQUENCE 36 AA; 4311 MW; E2A32293A866611C
                                                                                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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PRINTS; PR00278; PANCHORMONE. ProDom: PD001267; Pancreatic_hormn; 1. SMART; SM00309; PAH; 1.
PROSITE: PS00265; PANCREATIC_HORMONE_1; PROSITE: PS50276; PANCREATIC_HORMONE_2;
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                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Intestine;
MEDLINE=92319697; PubMed=1620652;
                                                                                                                                                 Query Match 78.9%;
Best Local Similarity 72.2%;
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36 36
36 AA; 4279 MW;
                                                                                                                                                                                                                                                                                                                     STANDARD;
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P80167;
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P29204;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                      RESULT 14
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    DR DR DR DR DR DR SQ SQ SQ
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                                                                                                                     Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Paracanthopterygii; Gadiformes; Gadidae; Gadus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.4%; Score 152; DB 1; Length 36; 72.2%; Pred. No. 1.8e-14; ive 6; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          AMIDATION.
17B09AA83867A7B6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00265; PANCREATIC_HORMONE_1; 1. PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
                                                                                                                                                                                                                                                                                                         Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                          TISSUE=Brain;
MEDLINE=93092973; PubMed=1459125;
                                                                                                                                                                                                                                                                                                                                                                                                                                         36 AA; 4267 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation.
                                                                                                              Jensen J., Conlon J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
nes 26; Conserv
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SEQUENCE
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Search completed: February 27, 2003, 14:36:51 Job time : 30 secs

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February 27, 2003, 14:32:08 ; Search time 29 Seconds
(without alignments)
255.783 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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194
1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                 671580 seqs, 206047115 residues
                                                                                                                                                                                                                                                                                                                           Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                     OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                  sp_vertebrate:*
sp_unclassified:*
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sp_bacteriap:*
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Maximum DB seq length: 2000000000
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Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Q9tr93 oryctolagus	Q91xd0 mus musculu	Q9tr92 oryctolagus	090wf4 paralichthy	Q919d3 ictalurus p	Q9dqk7 cyprinus ca	Q9n0m5 sus scrofa	Q9xsw6 macaca mula	Q925v2 mus musculu	Q9tsi6 ovis aries	Q8spf7 ovis aries	Q9pw68 typhlonecte	Q90wf3 paralichthy	Q9ps46 scyliorhinu	Q90wf2 paralichthy	Q9gk10 sūs scrofa
SUMMARIES	QI	Q9TR93	Q91XD0	Q9TR92	Q90WF4	Q919D3	Q9DGK7	O9N0M5	09XSW6	Q925V2	Q9TSI6	Q8SPF7	Q9PW68	Q90WF3	Q9PS46	Q90WF2	Q9GKL0
	DB	9	11	9	13	13	13	9	9	11	9	9	13	13	13	13	9
	Query Match Length DB	36	98	34	66	95	96	97	97	89	90	97	97	66	36	97	59
а₽	Query	96.9	94.8	90.7	77.8	76.8	75.8	75.3	74.2	73.7	73.7	73.7	72.7	72.7	70.1	68.0	50.5
	Score	188	184	176	151	149	147	146	144	143	143	143	141	141	136	132	98
	Result No.	7	7	m	4	S	9	7	80	6	10	11	12	13	14	15	16

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Gaps

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Length 36; 1; Indels

Score 188; DB 6; Pred. No. 2.3e-18; 0; Mismatches 1;

Query Match 96.9%; Best Local Similarity 97.2%; Matches 35; Conservative

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SEQUENCE 36 AA; 4285 MW; 02D499C8086DCC8D CRC64;

PROSITE; PS00265; PANCREATIC_HORMONE_1; 1. PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.

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O9ps51 lampetra fl O9nri6 homo sapien O9urg9 lymnaea sta O27441 aplysia cal O8qmy3 cowpox viru O97492 streptococc	ਰਾਹ≻÷	Ogy5h6 drosophila Ogy5h6 drosophila Og6qu6 homo sapien Ogwul4 homo sapien	mus aero omo	omo aeno aegi	Q96c69 homo sapien Q54307 streptomyce Q9dbv3 mus musculu Q14160 homo earien	Ω	S		AA.	ed) sequence update) annotation update)	Vartakrata: Butalanetomi:	a g		ely J., Eysselein V.E.,	e YY, PYY(1-36) and PYY(3-36),	/ PYY FAMILY.			,
Q9PS51 Q9NR16 Q9U0S9 Q27441 Q8QMY3 Q97Q92	09011/ 090113 090114 091218	29/24/39 29/246 29/246 29/21/4 29/21/3	Q90wx4 Q91XX1 Q9XD78 Q9NW90	Q96GG1 O16850 Q8S3W0	Q96C69 Q54307 Q9DBV3	Q8WWV8 Q9XH20	ALIGNMENT		PRT; 36	Created) Last sequenc Last annotat	t).	Lagomorpha; Leporidae; Oryctol	.000	ruk K., Shively	ms of peptide	HE NPY / PPY	Pancreatic_hormn.		Pancreatic_hormn; 1. AH; 1.
13 5 12 16	3 8 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	ነ ሊ ፋ ፋ ፋ ‹	11 17 4	4 10	111				ξ¥;	13,	(Rabbit) hordata:	gomorp	200 - 100	M., St	wo for	94).	ancrea	3; 1.	reation
10 10 10 # N O O	.1 504 .1 734 .1 6420 .1 356								PRELIMINARY	(TrEMBLrel (TrEMBLrel (TrEMBLrel	PYY(1-36). cuniculus	ria; La		miczek	on of t	; -820(19 BELONG	RON. 1955; E	hormone PANCE	7; Panc
3 3 3 5 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	000000	27.	27.	27. 27. 27.	27. 27.	27.				0 (Tr 0 (Tr 11 (Tr	, PYY IS cun Meta	Euthe =9986	07573	Schi	, kee izati	5:815 RITY:	103; 1 IPR00	159;	000126
69 67 63 58 58 58 58	00 00 00 00 00 00 00 00 00 00 00 00 00	* .	53.5 53.5 53	53 53 53		53 52.5		1	JTR93	01-MAY-2000 01-MAY-2000 01-OCT-2001	eptide YY Cyctolagu	Mammalia; Eutheria; NCBI_TaxID=9986;	[1] SEQUENCE. MPDIINE=05075735. Duhmad=7004400.	randt D.,	Character	In the rabbit.:; Peptides 15:815-820(1994). -!- SIMILARITY: BELONGS TO THE NPY	HSSP; P01303; 1RON. InterPro; IPR001955;	fam; PFO(TINTS: PF	Protocol Francisco Protocol Francisco Protocol Francisco Particular SMART; SM00309; PAH; 1.
118 119 20 21 22 22	22 2 2 2 2 2 4 2 4 2 4 2 4 2 4 2 4 2 4	32 32 32 32	3 3 4 5 6 5 6	37 38 39	0 4 4 4 0 11 2 4	444		RESULT 09TR93	ID OS		OS Or		RN [1				DR HS	DR Pf	

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2 KPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 34

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Conservative
                                                PRELIMINARY;
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                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
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                                                                                                                                                                                                                                    NCBI_TaxID=8255;
                                                                                                                           Neuropeptide Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Neuropeptide Y.
                                                                                                                                                                                                                                                                                   TISSUE-BRAIN;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25;
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                                                               Q90WF4;
                                              090WF4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q919D3
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                RESULT 4
Q90WF4
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Goebell H., Reeve J.R.Jr.;
"Characterization of two forms of peptide YY, PYY(1-36) and PYY(3-36),
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide YY, PYY(3-36).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 90.7%; Score 176; DB 6; Length 34; Best Local Similarity 100.0%; Pred. No. 9.1e-17; Matches 33; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                               Strausberg R.;
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; BC010821; AAH10821.1; -
InterPro; IPR001955; Pancreatic_hormn.
Prodom; P0001267; Pancreatic_hormn; 1.
Prodom; P0001267; Pancreatic_hormn; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in the rabbit.";
Peptides 15:815-820(1994).
1- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
HSSP; P01303; IRON.
InterPro; IPR001955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                          PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
SEQUENCE 98 AA; 11064 MW; 7AF165A1052C3249 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34 AA; 4024 MW; 02D4E9C38BA5FC8D CRC64;
                                                       01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Unknown (protein for MGC:19143).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                            Score 184; DB 11; NO. 2.4e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29 YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY 64
                             98 AA.
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1; Mismatches
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Probom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
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                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 94.4%;
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                              PRELIMINARY;
                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                      NCBI_TaxID=10090;
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                                                                                                                                                                                                                     TISSUE-COLON;
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                           091XD0
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RESULT 2
Q91XD0
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Q9TR92
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Leonard J.B.K., Waldbieser G.C., Silverstein J.T.; "Neuropeptide Y (NPY) sequence and distribution in channel catfish (Ictalurus punctatus).";
                                                                                                                                                                                                                     Paralichthys olivaceus (Flounder).

Bukaryota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi;
Actinopterygii: Teleostei: Buteleostei: Neoteleostei;
Acanthomorpha: Acanthopterygii: Percomorpha: Pleuronectiformes;
Pleuronectoidei: Paralichthyidae: Paralichthys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kurokawa T., Suzuki T.; "Development of neuropeptide Y related peptides in the digestive organs during the larval stage of Japanese flounder, Paralichthys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77.8%; Score 151; DB 13; Length 99; 69.4%; Pred. No. 7.1e-13; Live 7; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AB055211; BAB63409.1; -.
InterPro; IPROJ1955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
ProDom; P0001267; Pancreatic_hormn; 1.
PROSITE; PS00265; PANCREATIC_HORNONE_1; UNKNOWN_1.
PROSITE; PS002076; PANCREATIC_HORNONE_2; 1.
SEQUENCE 99 AA; 11215 MW; GFEED47F24CF6498 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
EMBL; AF267164; AAF71617.1; -
                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2000 (TrEMBLrel. 15, Last sequence update) 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29 YPVKPENPGDDAPAEELAKYYSALRHYINLITRQRY 64
99 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PS00265; PANCREATIC_HORMONE_1; PROSITE; PS50276; PANCREATIC_HORMONE_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSSP; P01303; 1RON.
InterPro; IPR001955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ictalurus punctatus (Channel catfish)
                                                                   Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
   PRT;
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                                                            01-DEC-2001 (TrEMBLrel. 19, 01-DEC-2001 (TrEMBLrel. 19, 01-MAR-2002 (TrEMBLrel. 20,
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us-09-634-363-2.rspt

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74.28;
                                                                                                                                                                                                                                                                               75.3%;
                                                                                                                                                                                                                                       8596 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neuropeptide Y (Fragment).
                                                                                                                                                                                                                                                                                                                      25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY;
                                                                                                                                                             Amidation; Neuropeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cercopithecinae; Macaca
                                                                                                                                                                                                  >45
  HSSP; P01303; 1RON.
                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                       76 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neuropeptide Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amidation.
                                                                                                                                                                                                                     NON_TER
SEQUENCE
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                                                                                                                                                                             NON_TER
CHAIN
                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monkeys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0925V2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9MSX60
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Q925V2
                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 8
Q9XSW6
  δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yingwen L., Takeshi Y.;
Yingwen L., Takeshi Y.;
"Daily rhythmic gene expression of neuropeptide Y in discrete brain of common carp. Cyprinus carpio, under the condition of self feeding.";
Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
BEBBL; AF287347; AAG00549.1;
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HSSP; P01303; IRON.
InterPro; IPR001955; Pancreatic_hormn.
                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                  Cyprinus carpio (Common carp).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Cyprinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Proneuropeptide Y (Fragment).
Sus scrofa (Pig).
Bukaryota: Metazoa: Chordata; Craniata; Vertebrata: Buteleostomi;
                                                                                                 ;
0
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0
                                                       Score 149; DB 13; Length 95; Pred. No. 1.3e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 96;
                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
EMBL; AF264083; AAF72538.1; -.
                  9AE19EAFBE24C6B5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96 AA; 10987 MW; C6C5ABCD87688980 CRC64;
                                                                                                                                                                                                                                                                                                              01-WAR-2001 (TrEMBLrel. 16, Created)
01-WAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 147; DB 13;
Pred. No. 2.4e-12;
                                                                                                                                    1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                           1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76 AA
                                                                                               5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00265; PANCREATIC_HORMONE_1; 1. PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00159; hormone3; 1.
PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                95 AA; 10729 MW;
                                                       76.8%;
72.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.8%;
69.4%;
                                                   Query Match 76.8
Best Local Similarity 72.2
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best_Local Similarity 69.4
Matches 25; Conservative
                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
TISSUE=HYPOTHALAMUS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                              Neuropeptide Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matteri R.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=BRAIN;
Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                SEQUENCE
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Q9N0M5;
                                                                                                                                                                                                                                                                                               09DGK7;
                                                                                                                                                                                                                                                                          Q9DGK7
                                                                                                                                                                                                                                   RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
Abler L.A., Golos T.G., Terasawa E.;
"Developmental changes in NPY mRNA expression in female rhesus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                      Score 146; DB 6; Length 76;
Pred. No. 2.5e-12;
6; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 144; DB 6; Length 97;
Pred. No. 6.2e-12;
7; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
-1 - SIMITARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
EMBL; AFF162-280; AAD43583.1; -.
HSSP: P01303; 1RON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97 AA; 10840 MW; 2D2209BAC20BD5EE CRC64;
                                                                                                                                                                                                                                                                                                    84E40EC2A4F94B2C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRORY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10 YPSKPDNPGEDAPAEDLARYYSALRHYINLITRQRY 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29 YPSKPDNPGEDAPAEDMARYYSALRHYINLITRQRY 64
                                                                                                                                                                                                                                              NEUROPEPTIDE Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89 AA.
                                                                                                                                                                                                                                                                                                                                                                                                              6; Mismatches
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PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
InterPro; IPR001955; Pancreatic_hormn. Pfam; PF00159; hormone3; 1. PR10178; PR00278; PANCHORMONE. ProDom; PD001267; Pancreatic_hormn; 1. SAMRT; SM00309; PAH; 1. PROSTITE; PS00265; PANCREATIC_HORMONE_1; PROSTITE; PS50276; PANCREATIC_HORMONE_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PR00278; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Interpro; IPR001955; Pancreatic_hormn.
Pfam; PF00159; hormone3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
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Peptides 22:325-334(2001).

-! FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN SECRETION OF GONADOTROPHIN-RELEASE HORMONE (BY SIMILARITY).

-! SUBCELLULAR LOCATION: SECRETED (BY SIMILARITY).

-! TISSUE SPECIFICITY: EXPRESSED THROUGHOUT THE BRAIN WITH HIGHEST LEVELS OF EXPRESSION IN MEDIAL PALLIUM, BASAL FOREBRAIN, PREOPTIC AREA MIDBRAIN TEGMENTUM AND TRIGEMINAL NUCLEUS.

-! ARAS SPECTROMERY: WW-4246; MW-ERR-4; METHOD-MALDI.

-! SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.

HSSP: PO1303; 1RON.

InterPro; IPRO1955; Pancreatic_hormn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ebersole T.J., Conlon J.M., Goetz F.W., Boyd S.K.; "Characterization and distribution of neuropeptide Y in the brain of caecilian amphibian.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A., SEQUENCE OF 29-64, TISSUE SPECIFICITY, AND MASS
                                                                                                                            Ovis aries (Sheep).
Sukaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Gymnophiona; Caeciliidae; Typhlonectes.
NCBI_TaxID=8456;
                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=LLE DE FRANCE; TISSUE-HYPOTHALAMUS;
Pillon D., Bruneau G.;
"Nucleotide sequence of Ovine preproneuropeptide Y.";
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ417904; CAD10677.1;
                                                                                                                                                                                                                                                                                                                                                                                                                      6C2209A361CF8583 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                      Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 143; DB 6; I
Pred. No. 8.4e-12;
7; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
                                                                                                                                                                                                                                                                                                                                                                                  NEUROPEPTIDE Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Α
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Typhlonectes natans (Rio Cauca caecilian).
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                                                                                                                                                                                                                                                                                                                                                                   POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                       POTENTIAL
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ProDom; PD001267; Pancreatic_hormn: 1.
SMART; SM00309; PAH; 1.
                                       Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
   PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7;
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                                                                                                                                                                                                                                                                                                                                                                                                                        10750 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          73.78;
66.78;
                                  01-JUN-2002 (TrEMBLrel. 21, 01-JUN-2002 (TrEMBLrel. 21, 01-JUN-2002 (TrEMBLrel. 21,
                                                                                          Neuropeptide Y precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
 PRELIMINARY;
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64
97
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24; Conser
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68
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TISSUE=BRAIN;
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                  Q8SPF7
                                                                                                                                                                                                                                                                                                                                                                     SIGNAL
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                                                                                                                       MEDLINE-21077529; PubMed-11210195;
Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;
Traylor B.A., Wnek C., Schroeder D., Phillips S.J.;
"Multiple obesity OTLS identified in an intercross between the NZO (New Zealand obese) and the SM (small) mouse strains.";
Mamm. Genome 12:95-103(2001).
Mamm. Genome 12:95-103(2001).
InterPro; IPR001955; Pancreatic_hormn.
PF00159; hormone3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovis aries (Sheep).

Eukaryota: Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
NCBI_TaxID=9940;
               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                    Length 89;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Simmons J.M., Daniel J.A., Matteri R.L., Keisler D.H.;
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
--- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
EMBL; AF095781, AACG9886.1; --- HSSP; P01303; IRON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90
9916 MW; 46FF2FB153EE5FFB CRC64;
                                                                                                                                                                                                                                                                                                                                                SEQUENCE 89 AA; 9943 MW; AE6052615A59D96A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-WAY-2000 (TrEMBLrel. 13, Created)
01-WAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Neuropeptide Y (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 143; DB 6; Lendon No. 7.8e-12;
                                                                                                                                                                                                                                                                 ProDom; PD001267; Pancreatic_hormn; 1.
PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
PROSITE; PS50276; PANCREATIC_HORMONE_2; 1.
NON_TER 89 89
                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Rest Local Similarity 66.7%; Pred. No. 7.7e-12;
Matches 24; Conservative 7; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 YPIKPEAPGEDASPEELNRYYASIRHYLNLVTRQRY 36
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Best Local Similarity 66.7%;
Matches 24; Conservative
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musculus (Mouse)
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90 AA;
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                                                 NCBI_TaxID=10090;
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  on pair of basic residues; Signal; Amidation
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
Neuropeptide Y, NPY=PANCREATIC polypeptide homolog.
Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Bukaryota: Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes; Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Crániata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Neopterygii; Reteostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Paralichthyidae; Paralichthys.
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AMIDATION (G-65 PROVIDE AMIDE GROUP)
(BY SIMILARITY).
9682E9D89DFFBLC7 CRC64;
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                                                                                                          97;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          olivaceus.";
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AB055212; BAB62410.1;
InterPro; IPR001955; Pancreatic_hormn.
Pfoon: PD001267; Pancreatic_hormn: 1.
ProDom: PD001267; Pancreatic_HORMONE_1; UNKNOWN_1.
PROSITE; PS002076; PANCREATIC_HORMONE_2; 1.
SEQUENCE 99 AA: 11179 MW; 32F6C21217CB1984 CRC64;
                                                                                                                                    Indels
                                                                                                          Length
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Last annotation update)
                                                                                                         Score 141; DB 13;
Pred. No. 1.6e-11;
                           NEUROPEPTIDE Y.
                                                                                                                                                          1 YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY 36
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                                                                                11275 MW;
                                                                                                         72.78;
63.98;
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                                                                                                                                23; Conservative
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Best Local Similarity
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                                                                             97 AA;
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68
68
 Neuropeptide;
SIGNAL
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Conlon J.M., Bjenning C., Hazon N.; "Structural characterization of neuropeptide Y from the brain of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Paralichthys olivaceus (Flounder).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Paralichthyidae; Paralichthys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kurokawa T., Suzuki T.;
"Development of neuropeptide Y related peptides in the digestive
organs during the larval stage of Japanese flounder, Paralichthys
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Pred. No. 2.6e-10;
8; Mismatches 6; Indels
                                                                                                                                                                                                                                         Length 36;
                                                                                                                                                                                                                                                                       Indels
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                          dogfish, Scyliorhinus canicula.";
Peptides 13:493-497(1992).
-!- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam: PF00159; hormone3; 1.
ProDom: PD001267; Pancreatic_hormn: 1.
PROSITE; PS00265; PANCREATIC_HORMONE_1; UNKNOWN_1.
SEQUENCE PS50776; PANCREATIC_HORMONE_2; 1.
SEQUENCE 97 AA; 10929 MW; 8703D0D393EF1E75 CRC64;
                                                                                                                                                                                                            36 AA; 4169 MW; 0D1715D0D9BD0DD6 CRC64;
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Last annotation update)
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Pred. No. 2.5e-11;
                                                                    HSSP; P01303; IRON.
InterPro; IPR001955; Pancreatic_hormn.
Pfam; PF00159; hormnoms; 1.
PRINTS; PR00128; PANCHORMONE.
ProDom; PD001267; Pancreatic_hormn; 1.
SMART; SM00309; PAH; 1.
PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.
PROSITE; PS00265; PANCREATIC_HORMONE_2; 1.
Amidation.
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22; Conservative
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Best Local Similarity
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01-MAR-2002
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